CTL in cancer patients. The present sequence represents a peptide from

the present invention

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The present invention describes peptides recognising cancer cells with tumour specific cytotoxic T lymphocytes (CTL). The peptides recognising cancer cells with tumour specific CTLs are selected from: (1) peptides of sequences (AAG68066 to AAG68069); (2) peptides containing the above mentioned sequences; (3) peptides having 70 % or more of homogeneity with the above mentioned sequences; and (4) peptides with one or more deleted, substituted, added or inserted amino acid(s) of the above mentioned sequences, particularly those having recognising property due to HLA-A2402 binding CTL, especially having at least 5 amino acids, used for medicine, particularly anticancer agents, derived from antitumour antigenic peptides of lck, src family, SART-1, SART-3 or cyclophilin B genes. The antitumour peptides have cytostatic activities. The peptides are used for the treatment of cancer. The peptides cause activation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 8; Page 2; 14pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JP2001245675-A.
                                                                                                                                                                                                        Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ITOY/) ITO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                     AAG68089;
                                                                                                                                                                                                                                                                                                                                                   RESULT 21
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This invention describes the novel use of interferons (IFNs) or DNAs capable of expressing the interferons and/or antigenic proteins (AP), antigenic peptides derived from the proteins or DNAs capable of expressing the antigenic proteins or peptides, in the manufacture of an agent for induction of antigen-specific T cells. The products of the invention have virucide and cytostatic activity and can be used for gene therapy or as inducers of antigen-specific T cells. The action of interferon-alpha (IFN-alpha) in a system for inducing specific cytotoxic T cell (CTL) by administering an antigenic peptide in an incomplete Freund's adjuvant (IFA) emulsion preparation form was evaluated. IFNs (or DNA encoding IFNs) are useful in the manufacture of a medicament for inducing antigen-specific T cells in an individual who has been administered with AP (or DNA encoding AP) or vice versa. The medicament is is useful for the treatment or prophylaxis of a tumor or a viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New peptides for recognizing cancer cells with tumor specific cytotoxic Tlymphocytes and for treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antitumour; cancer; cancer cell recognition; antigenic; CTL; lck; src; tumour specific cytotoxic T lymphocyte; anticancer; SART-1; SART-3; cyclophilin B gene; HLA-A2402.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.6%; Score 50; DB 4; Length 9; 100.0%; Pred. No. 1.4e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Prec. ....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antitumour peptide cyclophilin B 91-99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAG68089 standard; peptide; 9 AA.
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Best Local Similarity 100.
Matches 9; Conservative
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Gaps

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The invention relates to a composition (I) comprising an immune effector cell and a cell penetrating peptide (CPP) associated with an antigen or antibody. Also included are (I) a vaccine comprising (I), CPP associated with an antigen, and a pharmaceutically acceptable carrier and (2) corprehensing a composition for a disease, by providing (I) and CPP corprehensing a composition for disease, and introducing the antigens associated CPP to (I), where antigen enters into the cell. The antigens corprehensive (CC are, for example, tumour antigen derived epitopes recognised by tumour corprehension is useful for enhancing immunity in an animal to a disease, by administering a mature dendritic cell comprising CPP cor II. The composition is useful for enhancing immunity in an animal to a disease, by administering a mature dendritic cell comprising CPP corprehensive both CD4+ and CD8+ T cells. It is also useful for treating a comcer, non-Hodgkin's lymphoma, leukaemia, Hodgkin's lymphoma, uterine cancer, non-Hodgkin's lymphoma, leukaemia, Hodgkin's lymphoma, uterine cancer, cervical cancer, ovarian cancer and pancreatic cancer).

The animal is further subjected to a cancer treatment including surgery,
                                                                                                                                  0;
                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                        Cell penetrating peptide; cancer; tumour; melanoma; thymoma; antigen; lymphoma; sarcoma; lung cancer; non-Hodgkin's lymphoma; leukaemia; Hodgkin's lymphoma; uterine cancer; cervical cancer; bladder cancer; kidney cancer; adenocarcinoma; breast cancer; prostate cancer; ovarian cancer; pancreatic cancer; epitope; vaccine; dendritic cell; tumour infiltrating lymphocyte; TIL; human leukocyte antigen; HLA; cytostatic; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel composition for treating a disease in an animal, comprises an immune effector cell and cell penetrating peptide associated with an
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                                                                                                                               0; Indels
                                                                                            Length 9;
                                                                                        4.6%; Score 50; DB 4; Le
100.0%; Pred. No. 1.4e+06;
tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                        ABG79076 standard; peptide, 9 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-FEB-2002; 2002WO-US005212
                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                      Query Match
Best Local Similarity 100.0
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                                                    Sequence 9 AA;
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                                                                                                                                                                                                                                                                                                                                                                    15-NOV-2002
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ID ABG7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antigen specific T-cell; detection; diagnosis; cancer specific T-cell; cancer; tumour; cervical cancer; prostate cancer; cellular immunity; immune therapy; cytostatic; immunostimulant; vaccine; antigenic peptide; human; human leukocyte antigen; HIA-A24 epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 leukocyte antigen) peptides of human origin used in an example from the
radiation, chemotherapy or gene therapy. The administration of (I), preferably dendritic cell is prior to, subsequent to or concurrent with, the cancer treatment. The present sequence is a tumour antigen derived epitope for inclusion in the composition of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A detection method of antigen specific T-cells, comprises the use of plural antigenic peptides, useful in semi-quantitative determination of cancer specific T-cell frequencies and for monitoring cellular immunity.
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                                                                                                                                                                                    0; Indels
                                                                                                                                               Length 9;
                                                                                                                                               DB 5; Le
1.4e+06;
                                                                                                                                               4.6%; Score 50; DB 100.0%; Pred. No. 1.4 iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human CypB HLA-A24 epitope, SEQ ID NO:24.
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                                                                                                                                                                                                                                                                                                                                                                      ABR84374 standard; peptide; 9 AA.
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Best Local Similarity luv..
-Lac 9; Conservative
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                                                                                                                                                                                        9; Conservative
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Best Local Similarity
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                                                                                                          Sequence 9 AA;
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The invention relates to allergic reaction suppressors, comprising a peptide which originates from the same antigenic substance as the antigenic substance inducing the (peptide-originated) allergic reaction, and containing an epitope different from the epitope participating in the induction of the allergic reaction and yet dos not induce the allergic reaction. The desensitizers or allergic reaction suppressors are useful for treating and preventing type I allergic diseases and in cancer vaccines for preventing or treating cancer. The allergic reaction suppressors are also useful for suppressing an allergic reaction or reducing immunoglobulin E antibody production, and for desensitization to antigens. This sequence represents a peptide used in the invention and corresponds to amino acids 91-99 of the cyclophilin B protein.
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                                                                                                                                                                                 cytostatic; antiallergic; vaccine; allergic reaction suppressor; antigen; allergy; epitope; cancer; immunoglobulin E antibody; desensitization; cyclophilin B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Desensitizers or allergic reaction suppressors containing peptides originating from the same antigenic substances as reaction-inducing substances, useful for preventing or treating type I allergic diseases
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100.0%; Pred. No. 1.4e+06;
iive 0; Mismatches 0;
                                                                                                                                                             Cyclophilin B protein amino acids 91-99.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA.
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                                                                         ADC17702 standard; peptide; 9
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DEMIQGEDE 9
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                                                                                                                                                                                                                                                                                     WO2003020306-A1.
                                                                                                                                                                                                                                                                                                                                                                                                          (ITOH/) ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 9 AA;
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                                                                                                                                                                                                                                                                                                                  13-MAR-2003,
                                                                                                                                                                                                                                                       Synthetic.
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                                                                                                      ADC17702;
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4.6%; Score 50; DB 6; Length 9; 100.0%; Pred. No. 1.4e+06; tive 0; Mismatches 0; Indels

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This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                  This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
     Tumor antigen peptides derived from cyclophilin B for treatment and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
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100.0%; Pred. No. 1.7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                  Length 11;
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Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 11; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human cyclophilin B peptide fragment #18.
                                                                                                                                                                                                                                                                                                                                                                                AA.
                                                   Claim 4; Page 53; 64pp; Japanese
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                                                                                                                                                                                                                                                                                                                                                                             AAY69938 standard; peptide; 11
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Best Local Similarity 100..
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                                                                                                                                                                                                                                                                  8 ALIAGSVFFLL
                      diagnosis of tumors.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
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                              Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                        Tumor antigen peptides derived from cyclophilin B for treatment and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.6%; Score 50; DB 3; Length 10; 100.0%; Pred. No. 1.5e+02; cive 0; Mismatches 0; Indels
Human cyclophilin B peptide fragment #19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human cyclophilin B peptide fragment #15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA.
                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; Page 54; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY69935 standard; peptide; 11
                                                                                                                                                                                                                                          (SUMU ) SUMITOMO PHARM CO LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SUMU ) SUMITOMO PHARM CO LID. (ITOH/) ITOH K.
                                                                                                                                                                             99WO-JP003360
                                                                                                                                                                                                            98JP-00178449
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Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    42
                                                                                                                                                                                                                                                                                                                         WPI; 2000-116932/10.
                                                                                                                                                                                                                                                                                                                                                                         diagnosis of tumors.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-116932/10
                                                                                                                                                                                                                                                                                         Gomi S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ctoh K, Gomi S;
                                                                                                                                                                                                                                                          (ITOH/) ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cyclophilin B;
                                                                                                                                                                           24-JUN-1999;
                                                                                                                                                                                                           25-JUN-1998;
                                                                              Homo sapiens
                                                                                                           WO9967288-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9967288-A1
                                                                                                                                            29-DEC-1999
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                                                                                                                                                                                                                                                                                         Itoh K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 26
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Claim 8; Page 2; 14pp; Japanese.
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                        Gotoh M,
                                     WPI; 2001-193144/20
                                                                                                                                                                                                                                                                                  84 KFHRVIKDF
                                                                                                                                                                                                                                                                                               1 KFHRVIKDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                     JP2001245675-A
                                                                                                                                                                                                                                   Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ITOY/) ITO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                       Takasu H,
                                                                                                                                                                                                                                                                                                                                                        AAG68088;
                                                                                                                                                                                                                                                                                                                        RESULT 30
AAG68088
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                                                                                                                                                                                                                                                                         This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                      Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                               Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tumor antigenic peptide; interferon; IFN; antigen-specific T cell; virucide; cytostatic; gene therapy; IFN-alpha; cytotoxic T cell; viral infectious disease; cyclophilin B; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human cyclophilin B derived tumor antigenic peptide SEQ ID 16.
                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                 DB 3; Length 9;
1.4e+06;
                                                                                                                                                                                                                                                                                                                                4.5%; Score 49; DB 100.0%; Pred. No. 1.4 :ive 0; Mismatches
                                                        Human cyclophilin B peptide fragment #1
                                                                                                                                                                                                                                                           Claim 4; Page 49; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                        AAB46930 standard; peptide; 9 AA.
       AAY69921 standard; peptide; 9 AA
                                                                                                                                                                            (SUMU ) SUMITOMO PHARM CO LID (ITOH/) ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99JP-00207687
                                                                                                                                                             98JP-00178449.
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                                                                                                                                             99WO-JP003360.
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Best Local Similarity 100...
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                                                                                                                                                                                                                   WPI; 2000-116932/10.
                                                                                                                                                                                                                                                                                                                                                                 84 KFHRVIKDF
                                                                                                                                                                                                     Gomi S;
                                                                                                                                                                                                                                                                                                                  Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EP1074267-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                               Homo sapiens.
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                                                                                                                                             24-JUN-1999;
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                                        11-APR-2000
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                                                                                                                              29-DEC-1999.
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                         AAY69921;
                                                                                                                                                                                                    Itoh K,
                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 29
 AAB4693
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This invention describes the novel use of interferons (IFNs) or DNAs capable of antigenic peptides derived from the proteins or DNAs capable of expressing the antigenic proteins or DNAs capable of expressing the antigenic proteins or Deptides, in the manufacture of an agent for induction of antigen-specific T cells. The products of the invention have virucide and cytostatic activity and can be used for gene therapy or as inducers of antigen-specific T cells. The action of interferon-alpha (IFN-alpha) in a system for inducing specific cytotoxic crell (CTL) by administering an antigenic peptide in an incomplete or recell (CTL) by administering an antigenic peptide in an incomplete or administered with AP (or DNA encoding AP) or vice versa. The medicament con infectious disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New peptides for recognizing cancer cells with tumor specific cytotoxic T lymphocytes and for treating cancer.
                                                                                                                                                                                                                                     Use of antigenic proteins, peptides, interferon or their encoding DNA, in the manufacture of an agent for the induction of antigen-specific \ensuremath{\mathtt{T}}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antitumour; cancer; cancer cell recognition; antigenic; CTL; lck; src; tumour specific cytotoxic T lymphocyte; anticancer; SART-1; SART-3; cyclophilin B gene; HLA-A2402.
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 4.5%; Score 49; DB 4; Length 9; Best Local Similarity 100.0%; Pred. No. 1.4e+06; Matches 9; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 15; 25pp; English.
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(SUMU ) SUMITOMO PHARM CO LTD.
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The present invention describes peptides recognising cancer cells with tumour specific CTLB are selected from: (1) peptides of cancer cells with tumour specific CTLB are selected from: (1) peptides of sequences (AAG68066 to AAG68069); (2) peptides containing the above mentioned sequences; (3) peptides having 70 % or more of homogeneity with the above mentioned sequences; and (4) peptides with one or more deleted, substituted, added or inserted amino acid(s) of the above mentioned sequences, particularly those having recognising property due to HLA-A2402 binding CTL, especially having at least 5 amino acids, used for medicine, particularly anticancer agents, derived from antitumour medicine agents, derived from antitumour peptides have cytostatic activities. The peptides are used for the treatment of cancer. The peptides cause activation of CTL in cancer patients. The present sequence represents a peptide from the present invention Sequence 9 AA;

4.5%; Score 49; DB 4; Length 9; 100.0%; Pred. No. 1.4e+06; ive 0; Mismatches 0; Indels 100.0%; Preα. κ... Query Match Best Local Similarity luv... 9; Conservative 92 84 KFHRVIKDF ð

RESULT 31

ABG79075 standard; peptide; 9 AA. ABG79075; Human Cyp-B class I HLA widely expressed antigen peptide #1.

(first entry)

15-NOV-2002

penetrating peptide; cancer; tumour; melanoma; thymoma; antigen; lymphoma; sarcoma; lung cancer; non-Hodgkin's lymphoma; leukaemia; Hodgkin's lymphoma; uterine cancer; cervical cancer; bladder cancer; kidney cancer; adenocarcinoma; breast cancer; prostate cancer; ovarian cancer; pancreatic cancer; epitope; vaccine; dendritic cell; tumour infiltrating lymphocyte; TIL; human leukocyte antigen; HLA; cytostatic; human.

Homo sapiens.

WO200264057-A2.

22-AUG-2002

L5-FEB-2002; 2002WO-US005212.

15-FEB-2001; 2001US-0268687P

BAYU) BAYLOR COLLEGE MEDICINE

Wang R;

WPI; 2002-627577/67.

Novel composition for treating a disease in an animal, comprises an immune effector cell and cell penetrating peptide associated with an antigen or antibody.

Disclosure; Page 17; 61pp; English.

The invention relates to a composition (I) comprising an immune effector cell and a cell penetrating peptide (CPP) associated with an antigen or antibody. Also included are (1) a vaccine comprising (I), CPP associated with an antigen, and a pharmaceutically acceptable carrier and (2) preparing a composition for a disease, by providing (I) and CPP associated with an antigen for disease, and introducing the antigen-

infiltrating lymphocytes (TIL) of HLA (human leukocyte antigen) class I of infiltrating lymphocytes (TIL) of HLA (human leukocyte antigen) class I or II. The composition is useful for enhancing immunity in an animal to a disease, by administering a mature dendritic cell comprising CPP associated with an antigen to disease, to the animal, such that following the administration, animal is protected from disease, where the animal comprises both CD4+ and CD8+ T cells. It is also useful for treating a disease (e.g. cancer, tumour, melanoma, thymoma, lymphoma, arcoma, lung cancer, non-Hodgkin's lymphoma, leukaemia, Hodgkin's lymphoma, uterine cancer, cervical cancer, bladder cancer, kidney cancer, adenocarcinoma, breast cancer, prostate cancer, ovarian cancer and pancreatic cancer). The animal is further subjected to a cancer treatment including surgery, radiation, chemotherapy or gene therapy. The administration of (I), preferably dendritic cell is prior to, subsequent to or concurrent with, the cancer treatment. The present sequence is a tumour antigen derived epitope for inclusion in the composition of the invention associated CPP to (I), where antigen enters into the cell. The antigens

Sequence 9 AA;

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Gaps

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Gaps ·. 4.5%; Score 49; DB 5; Length 9; 100.0%; Pred. No. 1.4e+06; Live 0; Mismatches 0; Indels 0; Indels Query Match Best Local Similarity 100.

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92 84 KFHRVIKDF 1 KFHRVIKDF

à ద RESULT 32

ABR84373

ABR84373 standard; peptide; 9 AA. ABR84373;

(first entry) 06-NOV-2003

Human CypB HLA-A24 epitope, SEQ ID NO:23.

Antigen specific T-cell; detection; diagnosis; cancer specific T-cell; cancer; tumour; cervical cancer; prostate cancer; cellular immunity; immune therapy; cytostatic; immunostimulant; vaccine; antigenic peptide; human; human leukocyte antigen; HLA-A24 epitope.

Homo sapiens.

JP2002365286-A.

18-DEC-2002.

18-SEP-2001; 2001JP-00283413.

13-NOV-2000; 2000JP-00345094

(ITOY/) ITO Y.

WPI; 2003-508315/48.

A detection method of antigen specific T-cells, comprises the use of plural antigenic peptides, useful in semi-quantitative determination of cancer specific T-cell frequencies and for monitoring cellular immunity.

Example 8; Page 10; 18pp; Japanese.

The invention relates to a method for the detection of antigen specific T cells in a blood sample involving the use of a plurality of antigenic peptides. The method comprises sampling of peripheral blood monocytes; stimulation of the collected peripheral blood monocytes with antigens without direct use of antigen presenting cells; and detection of T-cells specific to the antigen in the stimulated monocytes. The method is particularly used for the detection of cancer as it can be used in semi-quantitative determination of cancer specific T-cells. It can also be used for cancer vaccine therapy for patients with cervical or prostate

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                                                                                                                                                                                                                                                                                                                                                                                                            Itoh K,
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                                                                   RESULT 34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to allergic reaction suppressors, comprising a peptide which originates from the same antigenic substance as the antigenic substance inducing the (peptide-originated) allergic reaction, and containing an epitope different from the epitope participating in the induction of the allergic reaction and yet dos not induce the allergic reaction. The desensitizers or allergic reaction suppressors are useful for treating and preventing type I allergic diseases and in cancer vaccines for preventing or treating cancer. The allergic reaction suppressors are also useful for suppressing an allergic reaction or reducing immunoglobulin E antibody production, and for desensitization to antigens. This sequence represents a peptide used in the invention and corresponds to amino acids 84-92 of the cyclophilin B protein.
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                                                                                                                                                                                                                                                                                                                                            cytostatic, antiallergic, vaccine, allergic reaction suppressor; antigen, allergy; epitope, cancer; immunoglobulin E antibody; desensitization; cyclophilin B.
cancer. The method can additionally be used to monitor of cellular immunity and cancer immune therapy by detection of specific T-cell frequencies. Sequences ABR84367-ABR84380 represent HLA-A24 (human leukocyte antigen) peptides of human origin used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Desensitizers or allergic reaction suppressors containing peptides originating from the same antigenic substances as reaction-inducing substances, useful for preventing or treating type I allergic diseases and in cancer vaccines.
                                                                                                                                    Gaps
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                                                                                                                                   0; Indels
                                                                                                          DB 6; Length 9;
1.4e+06;
                                                                                                          4.5%; Score 49; DB 100.0%; Pred. No. 1.4 iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                  Cyclophilin B protein amino acids 84-92.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim B; SEQ ID NO 1; 49pp; Japanese.
                                                                                                                                                                                                                                                      ADC17701 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-AUG-2002; 2002WO-JP008641
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                                                                                                                                     9; Conservative
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                                                                                                                                                                             84 KFHRVIKDF
                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Itoh K, Yamada A;
                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003020306-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ITOH/) ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9 AA;
                                                                                  Sequence 9 AA;
                                                                                                                                                                                                                                                                                                         18-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                          invention
                                                                                                                                                                                                                                                                               ADC17701;
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                                                                                                                                     Matches
                                                                                                                                                                                                                              RESULT 33
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This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
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                                                                                                                                                                                                                                                                    Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Escherichia coli; protein conformation; folding; acceleration; PPIase-alpha; peptidyl prolyl cis trans isomerase alpha; catalysis; isomerisation; prolyl peptide bond.
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100.0%; Pred. No. 1.4e+06;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                       Human cyclophilin B peptide fragment #30.
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                                                                                                                             AAY69950 standard; peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SUMU ) SUMITOMO PHARM CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98JP-00178449.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8; Conservative
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KFHRVIKDF 92
                                 Q)
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Best Local Similarity
                                   1 KFHRVIKDF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gomi
                                                                                                                                                                                                  11-APR-2000
                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                       24-JUN-1999;
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29-NOV-1995
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Gaps

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Katou S,
                                      WPI; 1995-140755/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diagnosis of tumors.
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  (TOFU ) TONEN CORP.
                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gomi S;
                                                                                                                                                                                              Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                          Cyclophilin B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9 AA
                                                                                                                                                                                                                                                                                                                                                      11-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9967288-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-DEC-1999.
                    Hayano T,
                                                                           proteins.
                                                                                                                                                                                                                                                                                                                                   AAY69963;
                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ITOH/)
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                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                RESULT
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                                                                                                                                                                                                                                                                                                                   o;
                                                                                                                                                                                                                    AAR72945-48 are tryptic fragments of E. coli PPIase-alpha (peptidyl prolyl cis trans isomerase). The E. coli PPIase-beta has a single mol. wt. of about 22 kDa and a single isoelectric point of about 9.7. The enzyme catalyses the isomerisation of prolyl peptide bonds in proteins and accelerates the folding of the protein. The inventors are claiming a PPIase-beta. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PR field.) (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                               New E.coli peptidyl prolyl cis trans isomerase beta - used to accelerate the folding of proteins, partic. for activation of inactive recombinant
                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Escherichia coli; protein conformation; folding; acceleration; PPIase-alpha; peptidyl prolyl cis trans isomerase alpha; catalysis; isomerisation; prolyl peptide bond.
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0
                                                                                                                                                                                                                                                                                                                       4.4%; Score 48; DB 2; Length 14; 76.9%; Pred. No. 3.9e+02; Live 0; Mismatches 3; Indels
                                                                                                                               Suzuki
                                                                                                                              Takahashi N,

 E. coli PPIase-alpha tryptic fragment 4.

                                                                                                                                                                                                                                                                                                                                                                                                                          AAR72903 standard; peptide; 14 AA.
                                                                                                                                                                                                     Example 7; Page 30; 85pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Escherichia coli; (ST 249 strain)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                89JP-00184738.
89JP-00260244.
89JP-00344705.
90EP-00307914.
                                                                      89JP-00260244.
                                             94EP-00203612
                                                                                                                              Maki N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94EP-00203610
                                                                                         90EP-00307914
                                                                                                                                                                                                                                                                                                                                                             88 VIKDFMIQGGDFT 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                              VIPGFMIQGGGFT 13
                                                                                                                                                                                                                                                                                                                                          10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
                                                                                                                             Katou S,
                                                                                                                                               WPI; 1995-140756/19
                                                                                                           (TOFU ) TONEN CORP.
                                                                                                                                                                                                                                                                                                                                Best_Local Similarity
Matches 10; Conser
                                                                                                                                                                                                                                                                                                       Sequence 14 AA;
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29-DEC-1989;
19-JUL-1990;
                                             19-JUL-1990;
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25-MAR-2003
29-NOV-1995
                                                                                          19-JUL-1990;
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29-DEC-1989
        EP647714-A1
                           12-APR-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EP647713-A1
                                                                                                                            Hayano T,
                                                                                                                                                                                  proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR72903;
                                                                                                                                                                                                                                                                                                                         Query Match
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AAR72903
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New E.coli peptidyl prolyl cis trans isomerase alpha - used to accelerate the folding of proteins, partic. for activation of inactive recombinant
                                                                                                                                                                                                                                                                                                                                                                                  AAR72900-04 are tryptic fragments of E. coli PPIase-alpha (peptidyl prolyl cis trans isomerase). The E. coli PPIase-beta has a single mol. wt. of about 22 kDa and a single isoelectric point of about 9.7. The enzyme catalyses the isomerisation of prolyl peptide bonds in proteins and accelerates the folding of the protein. The inventors are claiming the PPIase-alpha. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.) RR field.) (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tumor antigen peptides derived from cyclophilin B for treatment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.4%; Score 48; DB 2; Length 14; 76.9%; Pred. No. 3.9e+02; Live 0; Mismatches 3; Indels
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   Takahashi N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human cyclophilin B peptide fragment #43.
                                                                                                                                                                                                                                                                                                                  Example 7; Page 30; 85pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 10; Page 60; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY69963 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SUMU ) SUMITOMO PHARM CO LID
Maki N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-JP003360
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RESULT 39

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           capable of expressing the interferons and/or antigenic proteins (AP), antigenic peptides derived from the proteins or DNAs capable of expressing the antigenic proteins or peptides, in the manufacture of an agent for induction of antigen-specific T cells. The products of the invention have virucide and cytostatic activity and can be used for gene therapy or as inducers of antigen-specific T cells. The action of interferon-alpha (IFN-alpha) in a system for inducing specific cytotoxic T cell (CTL) by administering an antigenic peptide in an incomplete Freund's adjuvant (IFA) emulsion preparation form was evaluated. IFNS (or DNA encoding IFNS) are useful in the manufacture of a medicament for inducing antigen-specific T cells in an individual who has been administered with AP (or DNA encoding AP) or vice versa. The medicament is useful for the treatment or prophylaxis of a tumor or a viral infectious disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Use of antigenic proteins, peptides, interferon or their encoding DNA, the manufacture of an agent for the induction of antigen-specific T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes the novel use of interferons (IFNs) or DNAS
                                                                                                                                                                                                                                                                                                                      cell;
                                                                                                                                                                                                                                                                                                                      Tumor antigenic peptide; interferon; IFN; antigen-specific T cell virucide; cytostatic; gene therapy; IFN-alpha; cytotoxic T cell; viral infectious disease; cyclophilin B; human.
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                                                                                                                                                                                                                                                                                       Human cyclophilin B derived tumor antigenic peptide SEQ ID 33.
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                                  0; Indels
  Score 47; DB 3; Length 9; Pred. No. 1.4e+06;
                                   1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 20; 25pp; English.
                                                                                                                                                                                      AAB46947 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Takasu H, Gotoh M, Yamaoka T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SUMU ) SUMITOMO PHARM CO LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-JUL-2000; 2000EP-00306263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99JP-00207687.
  4.3%;
                                                                                                                                                                                                                                                       (first entry)
                                    8; Conservative
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                                                                    91 DFMIQGGDF 99
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Matches 8; Conser
Query Match
Best Local Similarity
                                                                                                  1 DYMIQGGDF
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                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                          EP1074267-A1
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                                                                                                                                                                                                                                                         04-MAY-2001
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                                                                                                                                                                                                                         AAB46947;
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                                   Matches
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The invention relates to allergic reaction suppressors, comprising a peptide which originates from the same antigenic substance as the antigenic substance inducing the (peptide-originated) allergic reaction, and containing an epitope different from the epitope participating in the induction of the allergic reaction and yet dos not induce the allergic reaction. The desensitizers or allergic reaction suppressors are useful for treating and preventing type I allergic diseases and in cancer vaccines for preventing or treating cancer. The allergic reaction suppressors are also useful for suppressing an allergic reaction or reducing immunoglobulin E antibody production, and for desensitization to antigens. This sequence represents a peptide used in the invention and corresponds to amino acids 91-99 of the cyclophilin B protein with the phe at position 92 replaced by a Tyr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ó
                                                                                                                                           cytostatic; antiallergic; vaccine; allergic reaction suppressor; antigen; allergy; epitope; cancer; immunoglobulin E antibody; desensitization; cyclophilin B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Desensitizers or allergic reaction suppressors containing peptides originating from the same antigenic substances as reaction-inducing substances, useful for preventing or treating type I allergic diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL;
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38.9%; Pred. No. 1.4e+06;
ive 1; Mismatches 0;
                                                                                                                 Modified cyclophilin B protein amino acids 91-99.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human cyclophilin B peptide fragment #42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 8; SEQ ID NO 9; 49pp; Japanese.
              ADC17709 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAY69962 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                 28-AUG-2002; 2002WO-JP008641.
                                                                                                                                                                                                                                                                                                                                                    29-AUG-2001; 2001JP-00260046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88.9%;
                                                                                (first entry)
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Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and in cancer vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9
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                                                                                                                                                                                                                                                                                                                                                                                                                       Yamada A;
                                                                                                                                                                                                                                                    WO2003020306-A1.
                                                                                                                                                                                                                                                                                                                                                                                      (ITOH/) ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-APR-2000
                                                                                18-DEC-2003
                                                                                                                                                                                                                                                                                    13-MAR-2003.
                                                                                                                                                                                                                   Synthetic.
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                                               ADC17709;
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ID AAY6
XX AAY6
AC AAY6
XX II-P
XX DE HUM3
XX CY01
ADC17709
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Claim 4; Page 55; 64pp; Japanese.
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                                                                                                                                                                                                                                          EP1074267-A1
                                                                                                                                                                                                                             sapiens
                                                                                                                                                                                                                                                                                 22-JUL-1999;
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                                                                                                                                                                                                                                                                                                             Takasu H,
                                                                                                                                                       AAB46946;
                                                               Query Match
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                                                                                                                           RESULT 42
AAB46946
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                                                                                                                                                                This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                 Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
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                                                                                                                                                                                                              Human cyclophilin B peptide fragment #21
HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                    Claim 10; Page 60; 64pp; Japanese.
                                                                                                                                                                                                                                                                                          AAY69941 standard; peptide; 9 AA.
                                                                                (SUMU ) SUMITOMO PHARM CO LID
                                                      99WO-JP003360.
                                                                                                                                                                                                                                                                                                                                                                                                                                         (SUMU ) SUMITOMO PHARM CO LID
                                                                   98JP-00178449
                                                                                                                                                                                                                 4.2%;
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                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                             92
                                                                                                                                                                                                                                                         Q.
                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                  WPI; 2000-116932/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-116932/10.
                                                                                                                                                                                                                                             84 KFHRVIKDF
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| KYHRVIKDF
                                                                                                      Gomi S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gomi S;
                                                                                       (ITOH/) ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ITOH K.
                                                                                                                                                                                                    Sequence 9 AA;
                                                      24-JUN-1999;
              Homo sapiens
                           WO9967288-A1
                                                                   25-JUN-1998;
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                                       29-DEC-1999
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                                                                                                     Itoh K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             toh K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ITOH/)
                                                                                                                                                                                                                                                                             RESULT 41
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This invention describes the novel use of interferons (IFNs) or DNAs capable of expressing the interferons and/or antigenic proteins (AP), antigenic peptides derived from the proteins or DNAs capable of expressing the antigenic proteins or peptides, in the manufacture of an agent for induction of antigen-specific T cells. The products of the invention have virucide and cytostatic activity and can be used for gene therapy or as inducers of antigen-specific T cells. The action of interferon-alpha (IFN-alpha) in a system for inducing specific cytotoxic T cell (CTL) by administering an antigenic peptide in an incomplete Freund's adjuvant (IFA) emulsion preparation form was evaluated. IFNs (or DNA encoding IFNs) are useful in the manufacture of a medicament for inducing antigen-specific T cells in an individual who has been administered with AP (or DNA encoding AP) or vice versa. The medicament is useful for the treatment or prophylaxis of a tumor or a viral
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This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tumor antigenic peptide; interferon; IFN; antigen-specific T cell; virucide; cytostatic; gene therapy; IFN-alpha; cytotoxic T cell; viral infectious disease; cyclophilin B; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Use of antigenic proteins, peptides, interferon or their encoding the manufacture of an agent for the induction of antigen-specific
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                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                  4.2%; Score 46; DB 3; Length 9; 100.0%; Pred. No. 1.4e+06; Live 0; Mismatches 0; Indels
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Length 9;

DB 4;

4.2%; Score 46;

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This sequence represents a cyclophilin B peptide of the invention. The
                                                                                                                                                                                                                              This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
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                                                                                                                                                                        Tumor antigen peptides derived from cyclophilin B for treatment diagnosis of tumors.
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                                                                                                                                                                                                                                                                                                          4.2%; Score 46; DB 3; Length 10; 100.0%; Pred. No. 4e+02; cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tumor antigen peptides derived from cyclophilin B for diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human cyclophilin B peptide fragment #13
                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY69933 standard; peptide; 11 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; Page 52; 64pp; Japanese.
                                                                                                                                                                                                            Claim 4; Page 53; 64pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SUMU ) SUMITOMO PHARM CO LID. (ITOH/) ITOH K.
                                                                                               (SUMU ) SUMITOMO PHARM CO LID. (ITOH/) ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98JP-00178449.
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                                                                                                                                  Gomi S;
                                                                                                                                                                                                                                                                                         Sequence 10 AA;
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           Gaps
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            Indels
 Pred. No. 1.4e+06;
; Mismatches 0;
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                                                                                                                                                                               Human cyclophilin B peptide fragment #17.
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                                                                                                               AAY69937 standard; peptide; 10 AA.
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             1;
                                                                                                                                                                                                                                                                                                                                                   (SUMU ) SUMITOMO PHARM CO LID. (ITOH/) ITOH K.
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88.9%;
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Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
                                                    KYHRVIKDF
 Best Local Similarity
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CC peptides are tumour antigen peptides derived from cyclophilin B, that CC recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The CC peptides are used for the treatment and diagnosis of tumours XX Squence 11 AA;
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peptides are used for the treatment and diagnosis of tumous Sequence 11 AA;

Query Match

Query Match

Best Local Similarity 100.0%; Pred. No. 4.6e+02;

Matches 11; Conservative 0; Mismatches 0; Indels

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Gaps

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Search completed: August 30, 2004, 10:49:40 Job time : 309.892 secs

4 LLAAALIAGSV 14 |||||||||| 1 LLAAALIAGSV 11

Sy da

us-09-720-469a-4.aug30.rag

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

August 30, 2004, 10:39:14; Search time 16.3885 Seconds (without alignments) 189.646 Million cell updates/sec Run on:

US-09-720-469A-4

1 GYKNSKFHRVI 11 Title: Perfect score:

Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

1586107 seqs, 282547505 residues Searched:

309569 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 8 Maximum DB seq length: 14

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

A_Geneseq_29Jan04:* 1: geneseqp1980s:*

geneseqp2001s:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	4 Human	Aay69921 Human cyc Aay69921 Human cyc	0 Human	8 Antitu	Abg79075 Human Cyp	3 Human		Aar72931 E. coli P	7 E.	Aay69962 Human cyc	Human	Aab27129 Human CAS	1	N	Abj15069 Human 125	Aay69959 Human cyc	Abr84746 Bovine po	Add88599 Influenza	Aaw19409 Anti-fund	Aaw19455 Anti-fung	5 Anti-	- 1	4 Anti-	7 Anti-
SO	AAY69924		AAB46930	AAG68088	ABG79075	ABR84373	ADC17701	AAR72931	AAR72887	AAY69962	AAB46946	AAB27129	AAW33943	AAW33942	ABJ15069	AAY69959	ABR84746	ADD88599	AAW19409	AAW19455	AAW19415	AAW19417	AAW19414	AAW19407
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Aaw19426 Anti-fung	Aaw19454 Anti-fung	Aaw19461 Anti-fung	Aaw19460 Anti-fung	Aaw19463 Anti-fung		Aay25141 Antibacte	Aay25143 Antibacte	Aay25140 Antibacte	Aab11990 Histatin-	Aab11976 Histatin-	Aab11985 Histatin-	Aab11987 Histatin-	Aab11977 Histatin-	Adb81288 Biologica	Aar89887 Human his	Aaw19441 Anti-fung	Aaw19452 Anti-fung	Aaw31675 Human his	Aaw49599 Human leu
AAW19426	AAW19454	AAW19461	AAW19460	AAW19463	AAW19472	AAY25141	AAY25143	AAY25140	AAB1199 0	AAB11976	AAB11985	AAB11987	AAB11977	ADB81288	AAR89887	AAW19441	AAW19452	AAW31675	AAW49599
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44.2	44.2	44.2	44.2	44.2	44.2	44.2	44.2	44.2	44.2	44.2	44.2	44.2		44.2		44.2	44.2	44.2	43.3
26.5	26.5	26.5	26.5	26.5	26.5	26.5	26.5	26.5	26.5	26.5	26.5	26.5		9		26.5	9	26.5	26
26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	4	42	43	44	45

ALIGNMENTS

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This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                              Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                              Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.0%; Score 60; DB 3; Length 11; Best Local Similarity 100.0%; Pred. No. 0.00013; Matches 11; Conservative 0; Mismatches 0; Indels
                                                                                                       Human cyclophilin B peptide fragment #4.
                          AAY69924 standard; peptide; 11 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; Page 50; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                    (SUMU ) SUMITOMO PHARM CO LID. (ITOH/) ITOH K.
                                                                                                                                                                                                                                                  99WO-JP003360.
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                                                                                                                                                                                                                                                                                                                                          Gomi S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 11 AA;
                                                                                                                                                                                               WO9967288-A1.
                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                  24-JUN-1999;
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                                                                                                                                                                                                                         29-DEC-1999.
                                                   AAY69924;
                                                                                                                                                                                                                                                                                                                                           Itoh K,
RESULT 1
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This invention describes the novel use of interferons (IFNS) or DNAs capable of expressing the interferons and/or antigenic proteins (AP), antigenic peptides derived from the proteins or DNAs capable of expressing the antigenic proteins or peptides, in the manufacture of an agent for induction of antigen-specific T cells. The products of the
                                                                                                                                                                                       This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                      Tumor antigen peptides derived from cyclophilin 3 for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumor antigenic peptide; interferon; IFN; antigen-specific T cell; virucide; cytostatic; gene therapy; IFN-alpha; cytotoxic T cell; viral infectious disease; cyclophilin B; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human cyclophilin B derived tumor antigenic peptide SEQ ID 16.
                                                                                                                                                                                                                                                                         53.3%; Score 32; DB 3; Lt
100.0%; Pred. No. 1.4e+06;
iive 0; Mismatches 0;
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                                                                                                                                                                    Claim 4; Page 49; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                             AAB46930 standard; peptide; 9 AA
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99WO-JP003360.
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                                               (SUMU ) SUMITOMO PHARM
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                                                           (ITOH/) ITOH K
                                                                                                                                                                                                                                                     Sequence 9 AA;
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24-JUN-1999;
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                                                                                                                                                         Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                              Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human cyclophilin B peptide fragment #1
                                                                                                                                    Human cyclophilin B peptide fragment #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY69921 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; Page 50; 64pp; Japanese
                                                             AAY69923 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                  (SUMU ) SUMITOMO PHARM CO LTD
                                                                                                                                                                                                                                                                                          98JP-00178449
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Best Local Similarity 100.v.
T; Conservative
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   GYKNSKFHRVI
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                                                                                                                                                                                                                                                                                                                                                     Gomi S;
                                                                                                                                                                                                                                                                                                                               ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 9 AA;
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                                                                                                                                                                                              Homo sapiens
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                                                                                      AAY69923;
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Gaps

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0; Indels Length 9;

DNA, in T

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invention have virucide and cytostatic activity and can be used for gene therapy or as inducers of antigen-specific T cells. The action of interferon-alpha (IFN-alpha) in a system for inducing specific cytotoxic T cell (CTL) by administering an antigenic peptide in an incomplete Freund's adjuvant (IFA) emulsion preparation form was evaluated. IFNs (or DNA encoding IFNs) are useful in the manufacture of a medicament for inducing antigen-specific T cells in an individual who has been administered with AP (or DNA encoding AP) or vice versa. The medicament is useful for the treatment or prophylaxis of a tumor or a viral
                                                                                                                                                                                                                                                                            infectious disease
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Sequence 9 AA;

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0; Indels
    Length 9;
53.3%; Score 32; DB 4; Le
100.0%; Pred. No. 1.4e+06;
iive 0; Mismatches 0;
                          6; Conservative
 Query Match
Best Local Similarity
                                                       6 KFHRVI 11
                                                                                 1 KFHRVI 6
               Best Loc
Matches
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RESULT 5

AAG68088 standard; peptide; 9 AA.

(first entry) 17-DEC-2001 AAG68088;

Antitumour peptide cyclophilin B 84-92.

Antitumour; cancer; cancer cell recognition; antigenic; CTL; lck; s; tumour specific cytotoxic T lymphocyte; anticancer; SART-1; SART-3; cyclophilin B gene; HLA-A2402.

Homo sapiens.

JP2001245675-A.

11-SEP-2001

25-DEC-2000; 2000JP-00393047

99JP-00374322 28-DEC-1999;

(ITOY/) ITO Y.

WPI; 2001-610076/70.

New peptides for recognizing cancer cells with tumor specific cytotoxic lymphocytes and for treating cancer.

Claim 8; Page 2; 14pp; Japanese.

The present invention describes peptides recognising cancer cells with tumour specific cytotoxic T lymphocytes (CTL). The peptides recognising cancer cells with tumour specific CTLs are selected from: (1) peptides of sequences (AAG68066 to AAG68069); (2) peptides containing the above mentioned sequences; (3) peptides having 70 % or more of homogeneity with the above mentioned sequences; and (4) peptides with one or more deleted, substituted, added or inserted amino acid(s) of the above mentioned sequences, particularly those having recognising property due to HLA-Sequences, particularly at least 5 amino acids, used for medicine, particularly anticancer agents, derived from antitumour cantigenic peptides of lck, src family, SART-1, SART-3 or cyclophilin B genes. The antitumour peptides have cytostatic activities. The peptides are used for the treatment of cancer. The peptides cause activation of cTL in cancer patients. The present sequence represents a peptide from the present invention

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Sequence 9 AA;
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 Score 32; DB 4; Length 9; Pred. No. 1.4e+06;
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Query Match
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6 KFHRVI 11
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RESULT 6 ABG79075

ABG79075 standard; peptide; 9 AA.

ABG79075;

(first entry) 15-NOV-2002

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Human Cyp-B class I HLA widely expressed antigen peptide #1.

Cell penetrating peptide; cancer; tumour; melanoma; thymoma; antigen; lymphoma; sarcoma; lung cancer; non-Hodgkin's lymphoma; leukaemia; Hodgkin's lymphoma; uterine cancer; cervical cancer; bladder cancer; kidney cancer; adenocarcinoma; breast cancer; prostate cancer; ovarian cancer; pancreatic cancer; epitope; vaccine; dendritic cell; tumour infiltrating lymphocyte; IIL; human leukocyte antigen; HLA; cytostatic; human.

Homo sapiens.

WO200264057-A2

22-AUG-2002

15-FEB-2002; 2002WO-US005212.

15-FEB-2001; 2001US-0268687P.

(BAYU) BAYLOR COLLEGE MEDICINE

Wang R;

WPI; 2002-627577/67.

Novel composition for treating a disease in an animal, comprises an immune effector cell and cell penetrating peptide associated with an antigen or antibody.

Disclosure; Page 17; 61pp; English.

The invention relates to a composition (I) comprising an immune effector cell and a cell penetrating peptide (CPP) associated with an antigen or antibody. Also included are (1) a vaccine comprising (I), CPP associated with an antigen, and apharmaceutically acceptable carrier and (2) preparing a composition for a disease, by providing (I) and CPP associated with an antigen for disease, and introducing the antigener cassociated with an antigen for disease, by providing (I) and CPP associated CPP to (I), where antigen enters into the cell. The antigens are, for example, tumour antigen for disease recognised by tumour infiltrating lymphocytes (TIL) of HLA (human leukocyte antigen) class I consist in the composition is useful for enhancing immunity in an animal to a disease, by administration, animal is protected from disease, where the animal consists both CD4+ and CD8+ T cells. It is also useful for treating a comprises both CD4+ and CD8+ T cells. It is also useful for treating a cancer, non-Hodgkin's lymphoma, leukaemia, thymoma, lymphoma, uterine cancer, non-Hodgkin's lymphoma, leukaemia, thymoma, ilmphoma, uterine cancer, cervical cancer, bladder cancer, kidney cancer, adenocarcinoma, breast cancer, prostate cancer, ovarian cancer and pancreatic cancer). The animal is further subjected to a cancer treatment including surgery, cradiation, chemotherapy or gene therapy. The administration of (I), the cancer treatment. The present sequence is a tumour antigen derived extreded extraction of the invention of the invention

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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                    Itoh K,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a method for the detection of antigen specific T cells in a blood sample involving the use of a plurality of antigenic peptides. The method comprises sampling of peripheral blood monocytes; stimulation of the collected peripheral blood monocytes with antigens without direct use of antigen presenting cells; and detection of T-cells specific to the antigen in the stimulated monocytes. The method is particularly used for the detection of cancer as it can be used in semi-used for cancer vaccine therapy for patients with cervical or prostate cancer. The method can additionally be used to monitor of cellular immunity and cancer immune therapy by detection of specific T-cell frequencies. Sequences ABR84367-ABR84380 represent HLA-A24 (human leukocyte antigen) peptides of human origin used in an example from the
                                                                                                                                                                                                                                                                                                                                                    Antigen specific T-cell; detection; diagnosis; cancer specific T-cell; cancer; tumour; cervical cancer; prostate cancer; cellular immunity; immune therapy; cytostatic; immunostimulant; vaccine; antigenic peptide; human; human leukocyte antigen; HIA-A24 epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A detection method of antigen specific T-cells, comprises the use of plural antigenic peptides, useful in semi-quantitative determination of cancer specific T-cell frequencies and for monitoring cellular immunity.
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                                                                    0; Indels
                                    53.3%; Score 32; DB 5; Length 9; 100.0%; Pred. No. 1.4e+06;
                                               100.0%; Preα. ....
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                      Human CypB HLA-A24 epitope, SEQ ID NO:23.
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                                                                                                                                                                                                                        ABR84373 standard; peptide; 9 AA
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                                                                                                      6 KFHRVI 11
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      Sequence 9 AA;
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                                     Query Match
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                                                                     Matches
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ABR84373
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The invention relates to allergic reaction suppressors, comprising a peptide which originates from the same antigenic substance as the antigenic substance inducing the (peptide-originated) allergic reaction, and containing an epitope different from the epitope participating in the induction of the allergic reaction and yet dos not induce the allergic reaction. The desensitizers or allergic reaction suppressors are useful for treating and preventing type I allergic diseases and in cancer vaccines for preventing or treating cancer. The allergic reaction suppressors are also useful for suppressing an allergic reaction or reducing immunoglobulin E antibody production, and for desensitization to antigens. This sequence represents a peptide used in the invention and corresponds to amino acids 84-92 of the cyclophilin B protein.
                                                                                                                                                                                             cytostatic; antiallergic; vaccine; allergic reaction suppressor; antigen; allergy; epitope; cancer; immunoglobulin E antibody; desensitization; cyclophilin B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Desensitizers or allergic reaction suppressors containing peptides originating from the same antigenic substances as reaction-inducing substances, useful for preventing or treating type I allergic diseases and in cancer vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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Pred. No. 1.4e+06;
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100.0%; Pred. No. 1.
                                                                                                                                                       Cyclophilin B protein amino acids 84-92.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 8; SEQ ID NO 1; 49pp; Japanese.
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ADC17701 standard; peptide; 9 AA.
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Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 9 AA;
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25-MAR-2003
29-NOV-1995
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AAR72886-92 are tryptic fragments of E. coli PPIase-beta (peptidyl prolyl cis trans isomerase). The E. coli PPIase-beta has a single mol. wt. of about 20 kDa and a single isoelectric point of about 5.0. The enzyme catalyses the isomerisation of prolyl peptide bonds in proteins and accelerates the folding of the protein. The inventors are claiming the PPIase-alpha. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PR field.) (indated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                            New E.coli peptidyl prolyl cis trans isomerase alpha - used to accelera the folding of proteins, partic. for activation of inactive recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 11;
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                                                                                                                                Suzuki
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                                                                                                                              Takahashi N,
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                                                                                                                                                                                                                                                                                            Example 6; Page 24; 85pp; English.
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    89JP-00260244.
89JP-00344705.
90EP-00307914.
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                                                                                                                              Maki N,
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nes 5; Conservative
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                                                                                                                              Katou S,
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YNNTIFHR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11 AA;
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                                                                                       (TOFU ) TONEN
      06-OCT-1989;
29-DEC-1989;
19-JUL-1990;
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                                                                                                                              Hayano T,
                                                                                                                                                                                                                                                     proteins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR72930-36 are tryptic fragments of E. coli PPIase-beta (peptidyl prolyl cis trans isomerase). The E. coli PPIase-beta has a single mol. wt. of about 20 kDa and a single isoelectric point of about 5.0. The enzyme catalyses the isomerisation of prolyl peptide bonds in proteins and accelerates the folding of the protein. The inventors are claiming a PPIase-beta. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PR field.) (Updated on 16-OCT-2003 to standardise OS field)
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PPIase-alpha; peptidyl prolyl cis trans isomerase alpha; catalysis; isomerisation; prolyl peptide bond.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Escherichia coli; protein conformation; folding; acceleration; PPIase-alpha; peptidyl prolyl cis trans isomerase alpha; catalysis; isomerisation; prolyl peptide bond.
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                                                                                                                                                                                                                                                                                                                                                                      Suzuki
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                                                                                                                                                                                                                                                                                                                                                                     Takahashi N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     coli PPIase-beta tryptic fragment 2.
                                                       Escherichia coli; (ST 249 strain).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 6; Page 24; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR72887 standard; peptide; 11 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Escherichia coli; (ST 249 strain)
                                                                                                                                                                                                                       89JP-00184738.
89JP-00260244.
89JP-00344705.
90EP-00307914.
                                                                                                                                                                                   94EP-00203612
                                                                                                                                                                                                                                                                                                                                                                     Maki N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94EP-00203610
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(first entry)
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Best Local Similarity
Matches 5; Conserv
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YNNTIFHR
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                                                                                                                                                                                                                                                                                                                           (TOFU ) TONEN
                                                                                                                                                                                                                                         06-OCT-1989;
29-DEC-1989;
19-JUL-1990;
                                                                                                                                                                                 19-JUL-1990;
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25-MAR-2003
29-NOV-1995
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                                                                                                                                          12-APR-1995
                                                                                                  EP647714-A1
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                                                                                                                                                                                                                                                                                                                                                                   Hayano T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         proteins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Use of antigenic proteins, peptides, interferon or their encoding DNA, in the manufacture of an agent for the induction of antigen-specific T
                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes the novel use of interferons (IFNs) or DNAs
recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                                                                              Tumor antigenic peptide; interferon, IFN; antigen-specific T cell; virucide; cytostatic; gene therapy, IFN-alpha; cytotoxic T cell; viral infectious disease; cyclophilin B; human.
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                                                                                                                                                                                                                                                                                                                                                                               Human cyclophilin B derived tumor antigenic peptide SEQ ID 32
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                                                                                  Score 29; DB 3; Length 9;
Pred. No. 1.4e+06;
.; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 19; 25pp; English.
                                                                                                                                                                                                                                                                             AAB46946 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yamaoka T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SUMU ) SUMITOMO PHARM CO LID
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Local Similarity 83.3%;
les 5; Conservative 1
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                                                                                     48.3%;
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                                                                    Query Match
Best Local Similarity 83.3
... 5; Conservative
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                                                                                                                                                        6 KFHRVI 11
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                                                                                                                                                                                        KYHRVI
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                                                     Sequence 9 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
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The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigenspecific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel CASB619 polypeptides useful for diagnosing, and as vaccines for prophylactic and therapeutic treatment of, cancers, particularly ovarian and colon carcinoma, and autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antagonist; integral membrane protein; inhibitor; transmembrane domain; therapy; psychotic disorder; Huntington's disease; Parkinson's disease; post-myocardial infarction; HIV infection; cell proliferation disorder; peptic ulcer; glaucoma; asthma; renal disease; autoimmune disease; antibacterial; betal-adrenergic receptor.
                                                                                                                                                                                                                       Human; CASB619; cancer; autoimmune disease; immunogen; vaccine; epitope.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Coche T, Vinals Y De BassolsC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2;
                                                                                                                                                                                          Human CASB619 protein epitope SEQ ID NO: 55.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                 (SMIK ) SMITHKLINE BEECHAM BIOLOGICALS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 7; Page 64; 68pp; English.
                                                                                               AAB27129 standard; protein; 10 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW33943 standard; peptide; 9 AA
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                                                                                                                                                                                                                                                                                                                                                                                    99GB-00007113
99GB-00022858
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|:||||
KYHRVI
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                                                                                                                                                                                                                                                                                         WO200058460-A2.
                                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                                               AAB27129;
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AAW33943
ID AAW33
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DE Betal
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KW Antag
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Query Match

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Human 125P5C8 epitope #3695
                                                                      Homo sapiens
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                                                                                                                                                                                                         27-MAR-1996;
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                                                                                                                                                                                                                                         20-AUG-1996;
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                                                                                                                                      02-OCT-1997
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(ODOW/)
                                                                                                                                                                                                                                                                            (NGGX/)
                                                                                                                                                                                                                                                                                              (SEEM/)
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ABJ15069
  BXXXXXB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence is a betal-adrenergic receptor antagonist, and is an antagonist of the invention. The antagonists (A) are for inhibiting the function of a prokaryotic or eukaryotic integral membrane protein (IMP) having at least one transmembrane domain (TMD), comprises a peptide including at least 4 consecutive amino acids (aa) from the sequence of the TMD. (A) are used to treat or prevent disorders in mammals that involve disturbances of IMP, and the same effect is achieved by inserting a nucleic acid encoding (A) into the calls of the mammal. Specified conditions that can be treated are schizophrenia, psychotic disorders, Huntington's disease, Tourette's syndrome, hypertension, post-myocardial infarction, tachyarrhythmia, human immunodeficiency virus infection, AIDS, cell proliferation (cancer, psoriasis or hyperkeratosis), and drug abuse. (A) can also be used to treat peptic ulcer, glaucoma, asthma, renal disease, Parkinson's disease, congestive heart failure, migraine, and drug antagonic in the cancer of the constitution of the cancer of the c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           depression, obesity, diabetic neuropathy/retinopathy, autoimmune diseases, graft rejection etc. (A) can be used as diuretics and antibacterials. (A) may also be used to determine function of orphan receptors and, when attached to a marker, for tissue imaging to localise/quantify specific receptors. The transgenic animals are used to test efficacy/toxicity of (A) and as models of diseases. (A) are very specific and selective, against the IMP from which they are derived, so
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antagonist; integral membrane protein; inhibitor; transmembrane domain; therapy; psychotic disorder; Huntington's disease; Parkinson's disease;
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                                                                                                                                                                                                                                                                                                                                                                              Inhibitor of integral membrane protein - used to treat e.g. schizophrenia, hypertension, viral infection, cancer etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 28; DB 2; Length 9;
Pred. No. 1.4e+06;
1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                            Odowd BF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Betal-adrenergic receptor antagonist.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 17; Page 94; 127pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW33942 standard; peptide; 14 AA
                                                                                                                                                                                                                                                                                                           George SR,
                                                                                                                                                  96US-0014306P.
96US-00670119.
96US-0024240P.
                                                                                                                97WO-CA000203
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Best Local Similarity
Matches .5; Conserv
                                                                                                                                                                                                                                                                                                           Seeman P,
                                                                                                                                                                                                                    NG G Y K.
SEEMAN P.
GEORGE S R
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9 AA;
               Homo sapiens,
                                                 WO9735881-A2
                                                                                                                 26-MAR-1997;
                                                                                                                                                   27-MAR-1996;
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                                                                                 02-OCT-1997
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(SEEM/)
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This sequence is a betal-adrenergic receptor antagonist, and is an antagonist of the invention. The antagonists (A) are for inhibiting the function of a prokaryotic or eukaryotic integral membrane protein (IMP) having at least one transmembrane domain (TMD), comprises a peptide including at least 4 consecutive amino acids (aa) from the sequence of including at least 4 consecutive amino acids (aa) from the sequence of including at least 4 consecutive amino acids (aa) from the sequence of including at least 4 consecutive amino acids (ab from the sequence of including at least 4 consecutive amino acids (ab from the sequence of including at least 4 consecutive amino the same effect is achieved by inserting a nucleic acid encoding (A) into the cells of the mammal. Specified conditions that can be treated are schizophrenia, psychotic disorders, and infarction, tachyarrhythmia, human immunodeficiency virus infection, alboring and subsection (cancer, psoriasis or hyperkeratosis), and drug abuse. (A) can also be used to treat peptic ulcer, glaucoma, asthma, crenal disease, parkinson's disease, congestive heart failure, migraine, diseases, graft rejection etc. (A) can be used as diuretics and antibacterials. (A) may also be used to determine function of orphan receptors and, when attached to a marker, for tissue imaging to localise/quantify specific receptors. The transgenic animals are used to test efficacy/loxicity of (A) and as models of diseases. (A) are very specific and selective, against the IMP from which they are derived, so should have few side effects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ó
post-myocardial infarction; HIV infection; cell proliferation disorder; peptic ulcer; glaucoma; asthma; renal disease; autoimmune disease; antibacterial; betal-adrenergic receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Inhibitor of integral membrane protein - used to treat e.g. schizophrenia, hypertension, viral infection, cancer etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            46.7%; Score 28; DB 2; Length 14; 62.5%; Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Odowd BF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 17; Page 94; 127pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABJ15069 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         George SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96US-0014306P.
96US-00670119.
96US-0024240P.
                                                                                                                                                                                                                                                                                                                                                                                                               97WO-CA000203.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6 GYANSAFN 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NG G Y K.
SEEMAN P.
GEORGE S R.
O'DOWD B F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1997-489566/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GYKNSKFH 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ng GYK, Seeman P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
Matches 5; Conserv
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Gomi S;

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Itoh K,
                                                                                                                                                             Matches
                                                                                                                                                                                                                          RESULT 18
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                                                                                                                                                                                                                                         The present invention relates to compositions comprising a substance that modulates the status of 125P5C8 or a molecule that is modulated by 125P5C8. The status of a cell that expresses 125P5C8 is modulated. The composition is useful for treating cancer, particularly prostate, bladder, kidney, colon, ovary or breast cancer. The 125P5C8 protein and/or a nucleotide sequence encoding the protein is useful for immunising a mammal against cancer. The present sequence is a 125P5C8 epitope shown in the exemplification of the invention
                                                                                                                                                                                               125P5C8 gene or a molecule that is modulated by 125P5C8, useful for treating or preventing cancer that expresses or over expresses 125P5C8.
                                                                                                                                            Ge W;
          cancer; cytostatic; breast cancer; prostate cancer; kidney cancer; colon cancer; ovarian cancer; epitope.
                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL;
HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                       composition comprising a substance that modulates the status of
                                                                                                                                                                                                                                                                                                                                                             .;
                                                                                                                                            Raitano AB,
                                                                                                                                                                                                                                                                                                                                           Score 28; DB 5; Length 14; Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                           1; Indels
                                                                                                                                            Afar DEH,
                                                                                                                                                                                                                                                                                                                                                             2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human cyclophilin B peptide fragment #39.
                                                                                                                                            Hubert RS, 1
Jakobovits A;
                                                                                                                                                                                                                           Disclosure; Page 216; 274pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY69959 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SUMU ) SUMITOMO PHARM CO LID (ITOH/) ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98JP-00178449
                                                                                         2002WO-US007855.
                                                                                                         14-MAR-2001; 2001US-00809638.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99WO-JP003360
                                                                                                                                                                                                                                                                                                                                           46.7%;
                                                                                                                                            PM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                    Morrison RK, Morrison K,
                                                                                                                                                                                                                                                                                                                                                             4; Conservative
                                                                                                                                           Challita-Eid
                                                                                                                                                                     WPI; 2002-713510/77.
                                                                                                                                                                                                                                                                                                                                                                                        |:|:||
4 YENTHFH 10
                                                                                                                          (AGEN-) AGENSYS INC
                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                             2 YKNSKFH 8
                   bladder cancer;
         Human; 125P5C8;
                                                                                                                                                                                                                                                                                                                           Sequence 14 AA;
                                                     WO200272785-A2
                                     Homo sapiens
                                                                                        13-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-JUN-1999;
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                                                                      19-SEP-2002
                                                                                                                                            Faris M,
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The present invention provides the protein sequence of a polypyrimidine tract protein isoform. Also provided are methods for modulating the stability of a CD154 RNA. The methods are useful for increasing the stability of a RNA, decreasing the stability of a RNA, preventing or treating allograft rejection in a subject, inhibiting CD40 activation in a subject, and identifying agents that modulate the level or activity of a polypyrimidine tract protein. Agents which modulate the activity of a polypyrimidine tract protein are useful to prevent or treat allograft rejection or inhibit CD40 activation. The present sequence is a nonapeptide conserved between the bovine and human polypyrimidine tract
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A new polypyrimidine tract protein isoform useful for finding agents which modulate its activity and to prevent or treat allograft rejection or CD40 activation.
                                                                                                                                                                                                                  This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cow; polypyrimidine tract protein; isoform; RNA stability regulation; immunosuppressive; allograft rejection; CD154; PTB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                         Tumor antigen peptides derived from cyclophilin B for treatment and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bovine polypyrimidine tract binding protein conserved nonapeptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 27; DB 3; Le
Pred. No. 1.4e+06;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                       45.0%; Scc...
100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABR84746 standard; protein; 9 AA.
                                                                                                                                                         Claim 4; Page 61; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-JAN-2002; 2002US-0349869P. 02-JAN-2003; 2003US-0437779P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-JAN-2003; 2003WO-US001623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (DART-) DARTMOUTH COLLEGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                             diagnosis of tumors.
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WPI; 2000-116932/10.
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les 5; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FHRVI
                                                                                                                                                                                                                                                                                                                                                                                   Sequence 9 AA;
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AAW19409;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UXBO-)
                                                                                                                                                                                                                                                                                       Key
Region
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AAW19455
ID AAW19
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AC AAW19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New Replikin peptides of Plasmodium falciparum or influenza virus, useful for diagnosing, preventing or treating influenza virus infection or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated Plasmodium falciparum peptide or an influenza virus peptide. The composition and methods are useful in diagnosing, preventing or treating influenza virus infection or malaria. Also disclosed is an influenza virus vaccine comprising at least one isolated Replikin present in the haemagglutinin protein of an emerging strain of influenza virus and a pharmaceutical carrier and/or adjuvant and a malaria vaccine comprising at least one isolated P. falciparum Replikin and a pharmaceutical carrier. The present sequence represents the amino acid sequence of a Replikin sequence.
                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                           vaccine; gene therapy; malaria; influenza; Replikin; haemagglutinin.
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                                                                             3; Indels
                                     Score 27; DB 7; Length 9; Pred. No. 1.4e+06; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; SEQ ID NO 266; 136pp; English.
                                                                                                                                                                                                                                                               ADD88599 standard; peptide; 11 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW19409 standard; peptide; 11 AA.
                                                                                                                                                                                                                                                                                                                                                                                    Influenza virus H3N2 Replikin #4.
                                    45.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-MAR-2001; 2001US-0278761P.
09-JUL-2001; 2001US-0303396P.
26-OCT-2001; 2001US-00984057.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-MAR-2002; 2002US-00105232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          45.0%;
ilarity 57.1%;
Conservative
                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bogoch ES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-852213/79.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (BOGO/) BOGOCH S.
(BOGO/) BOGOCH E S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 NSKFHRV 10
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Best Local Similarity
Matches 4; Conserv
                                                         Local Similarity
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                                                                                                                   2 YKNSKFHR
                                                                                                                                                          YGNSPLHR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Orthomyxoviridae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2003180328-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 11 AA;
Sequence 9 AA;
                                                                               .;
2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-SEP-2003
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                                                                                                                                                                                                                                                                                                    ADD88599;
                                       Query Match
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                                                            Best Loc
Matches
                                                                                                                                                                                                                    RESULT 19
ADD88599
ID ADD88
XX
DX 29-JA
XX
DE Influ
XX
VX
OCC
CC Also

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ID AAW1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      - useful esp for
                                                                                                                      Candida albicans; peridontitis; caries; tooth decay; oral infection; vaginal infection, urethral infection; mucosal infection; ear infection; respiratory infection; skin infection; ophthalmic infection; bacterial disease; Streptococcus mutans; Porphyromonas gingivalis; Clostridium histolyticum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               on
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Friden PM;

    1. 11
/note= "At least one amino acid must have a
configuration"

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                                                            Anti-fungal and anti-bacterial histatin-based peptide 117.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 26.5; DB 2; Length 1
Pred. No. 3e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anti-fungal and anti-bacterial histatin-based peptide(s) treating Candidal infections, periodontitis and caries.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Spacciapoli P,
                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Roberts FD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW19455 standard; peptide; 11 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     96WO-US009374.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95US-00481888
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75.0%;
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PERI-) PERIODONTIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oppenheim FG, Xu T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GYKR-KFH 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1997-052232/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GYKNSKFH 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 11 AA;
04-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9640768-A2
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                                                                                                                                                                                                                                                                                                                            Synthetic
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configuration"
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75.0%;
                                                                                                                                                                                                                                                                                                                                                                            96WO-US009374.
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             05-SEP-1997 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                              (PERI -) PERIODONTIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Xu T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GYKR-KFH 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 12 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oppenheim FG,
                                                                                                                                                                                                                                                                                                                                                                            07-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                              07-JUN-1995;
                                                                                                                                                                                                                                                                                                         WO9640768-A2
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                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a specifically claimed example of an anti-
-fungal and anti-bacterial D-amino acid histatin-based peptide 117, where
-fungal and anti-bacterial D-amino acid histatin-based peptide 117, where
at least one amino acid in the peptide is in the D-configuration. The
peptide is preferably modified by an acetyl or carbamyl addition at the N
-terminus and/or amidation at the C-terminus. The novel D-amino acid-
containing peptide, based on the naturally occurring histidine-rich human
compositions for the treatment of oral, vaginal, urethral, mucosal,
composition, averand ophthalmic fungal or bacterial infections. It
is particularly effective against local and systemic Candida albicans
infection, against oral bacterial diseases such as caries and
periodontitis (by inhibiting e.g. Streptococcus mutans and Porphyromonas
gingivalis) and against Clostridium histolyticum. The D-amino acid-
containing peptide has superior anti-fungal (especially anti-candidal)
and anti-bacterial activity, particularly on a weight basis, compared to
the natural L- amino acid forms of histatin-based peptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1;
                                                              Candida albicans; peridontitis; caries; tooth decay; oral infection; vaginal infection; urethral infection; mucosal infection; ear infection; respiratory infection; skin infection; ophthalmic infection; bacterial disease; Streptococcus mutans; Porphyromonas gingivalis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anti-fungal and anti-bacterial D-amino acid histatins and histatin-based peptide(s) - useful esp for treating Candidal infections, periodontitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The presence of D-residues also makes the peptides more resistant to degradation than corresponding L-amino acid versions
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H
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    .11
/note= "At least one amino acid must have
configuration"

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                                Anti-fungal and anti-bacterial D-amino acid peptide 117.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Spacciapoli P,
                                                                                                                                                                                                       Location/Qualifiers
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6; Conservative
                                                                                                                                      Clostridium histolyticum
                                                                                                                                                                                                                                                                                                                                                                                                                              (PERI-) PERIODONTIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oppenheim FG, Xu T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 11 AA;
                                                                                                                                                                                                                                                                                                                                                           07-JUN-1996;
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05-SEP-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and caries.
                                                                                                                                                                      Synthetic
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AAW19415
ID AAW19
XX
AC AAW19
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AAW19415 standard; peptide; 12 AA

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AAW19415

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a specifically claimed peptide sequence, peptide 113-F5, where at least one amino acid is in the D-configuration. The peptide preferably is modified by an acetyl or carbamyl addition at the N-terminus and/or amidation at the C-terminus. The peptide, based on the naturally occurring histidine-rich human histatins, have anti-fungal and anti-bacterial activity and are useful in compositions for the treatment of oral, vaginal, urethral, mucosal, respiratory, skin, ear and ophthalmic fungal or bacterial infections. It is particularly effective against local and systemic Candida albicans infection, against oral bacterial diseases such as caries and periodontitis (by inhibiting e.g.

    useful esp for

                                                                   Candida albicans; peridontitis; caries; tooth decay; oral infection; vaginal infection; urethral infection; mucosal infection; ear infection; respiratory infection; skin infection; ophthalmic infection; bacterial disease; Streptococcus mutans; Porphyromonas gingivalis; Clostridium histolyticum.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1. .12
/note= "At least one amino acid must have a D-
Anti-fungal and anti-bacterial histatin-based peptide 113-F5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Friden PM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-fungal and anti-bacterial histatin-based peptide(s) treating Candidal infections, periodontitis and caries.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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Pred. No. 3.3e+02;
); Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roberts FD,
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pertide 113-F4.5, where at least one amino acid is in the D-
configuration. The peptide preferably is modified by an acetyl or
carbamyl addition at the N-terminus and/or amidation at the C-terminus.

The peptide, based on the naturally occurring histidine-rich human
histatins, have anti-fungal and anti-bacterial activity and are useful in
compositions for the treatment of oral, vaginal, urethral, mucosal,
respiratory, skin, ear and ophthalmic fungal or bacterial infections. It
is particularly effective against local and systemic Candida albicans
infection, against oral bacterial diseases such as caries and
periodontitis (by inhibiting e.g. Streptococcus mutans and Porphyromonas
gingivalis) and against Clostridium histolyticum. The peptide has
superior anti-fungal (especially anti-Candidal) and anti-bacterial
activity, particularly on a weight basin, compared to the longer,
naturally occurring histatins. Peptides containing D-residues are also
more resistant to degradation than L-amino acid versions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Jal and anti-bacterial histatin-based peptide(s) - useful esp for Candidal infections, periodontitis and caries.
                                             Candida albicans; peridontitis; caries; tooth decay; oral infection; vaginal infection; urethral infection; mucosal infection; ear infection; respiratory infection; skin infection; ophthalmic infection; bacterial disease; Streptococcus mutans; Porphyromonas gingivalis; Clostridium histolyticum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a specifically claimed peptide sequence,
               Anti-fungal and anti-bacterial histatin-based peptide 113-F4.5.
                                                                                                                                                                                                       1. .12
/note= "At least one amino acid must have a D-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Anti-fungal and anti-bacterial histatin-based peptide 113-F4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 26.5; DB 2; Length 12;
Pred. No. 3.3e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Roberts FD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW19414 standard; peptide; 12 AA.
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                                                                                                                                                                                                                                         configuration
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                                                                                                                                                                                                                                                                                                                                                                         95US-00481888
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Best Local Similarity 75.0
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          PERIODONTIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Xu T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6 GYKR-KFH 12
                                                                                                                                                                                                                                                                                                                                                                                                                          UNIV BOSTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1997-052232/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 12 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oppenheim FG,
                                                                                                                                                                                                                                                                                                                                                                         07-JUN-1995;
                                                                                                                                                                                                                                                                       WO9640768-A2
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                                                                                                                                                                                                                                                                                                       19-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Anti-fungal
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                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              treating
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                                                                                                                                                                                                     Region
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                                                                                                                                                                                        Key
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EX BX BX B
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Friden PM;

Spacciapoli P,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anti-fungal and anti-bacterial histatin-based peptide(s) - useful esp for treating Candidal infections, periodontitis and caries.
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                             vaginal infection; urethral infection; mucosal infection; ear infection; respiratory infection; skin infection; ophthalmic infection; bacterial disease; Streptococcus mutans; Porphyromonas gingivalis; Clostridium histolyticum.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Streptococcus mutans and Porphyromonas gingivalis) and against
Clostridium histolyticum. The peptide has superior anti-fungal
(especially anti-Candidal) and anti-bacterial activity, particularly o
weight basin, compared to the longer, naturally occurring histatins.
Peptides containing D-residues are also more resistant to degradation
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             Candida albicans; peridontitis; caries; tooth decay; oral infection;
                                                                                                                                                              1. .12
/note= "At least one amino acid must have a D-
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                                                                                                                                                                                                                                                                                                                                                                                                            Friden PM;
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Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW19407 standard; peptide; 12 AA.
                                                                                                                                                                                               configuration"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; Fig 1; 72pp; English
                                                                                                                                                                                                                                                                                            96WO-US009374.
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75.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6; Conservative
                                                                                                                                                                                                                                                                                                                                                          PERIODONTIX INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12
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Matches 6; Conserv
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                                                                                                                                                                                                                                                                                          07-JUN-1996;
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                                                                                                             Synthetic.
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                                                                                                                                              Key
Region
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Candida albicans; peridontitis; caries; tooth decay; oral infection;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence represents a specifically claimed peptide sequence, peptide 113, where at least one amino acid is in the D-configuration. The peptide preferably is modified by an acetyl or carbamyl addition at the reminus and/or amidation at the C-terminus. The peptide, based on the naturally occurring histidine-rich human histatins, have anti-fungal and anti-bacterial activity and are useful in compositions for the treatment of oral, vaginal, urethral, mucosal, respiratory, skin, ear and ophthalmic fungal or bacterial infections. It is particularly effective against local and systemic Candida albicans infection, against oral bacterial diseases such as caries and periodontitis (by inhibiting e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      - useful esp for
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vaginal infection; urethral infection; mucosal infection; ear infection; respiratory infection; skin infection; ophthalmic infection; bacterial disease; Streptococcus mutans; Porphyromonas gingivalis; Clostridium histolyticum.
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                                                                                                                                                                                                                                                                                                                                                                                      Friden PM;
                                                                                                                                                    /note= "At least one amino acid must have a
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Pred. No. 3.3e+02;
0; Mismatches 1;
                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                        Roberts FD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW19426 standard; peptide; 12 AA
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                                                                                                                                                                       configuration"
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75.0%;
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                                                                                                                                                                                                                                                                                                                                                                                       Xu T,
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                                                                                                                                                                                                                                                                                                                                                      UNIV BOSTON
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Matches 6; Conserv
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                                                                                      Synthetic
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Region
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The present sequence represents a specifically claimed peptide sequence, peptide 113-Q3.9, where at least one amino acid is in the D-configuration. The peptide preferably is modified by an acetyl or carbamyl addition at the N-terminus and/or amidation at the C-terminus.

CC carbamyl addition at the N-terminus and/or amidation at the C-terminus.

CC carbamyl addition at the N-terminus and/or amidation at the C-terminus.

CC The peptide, based on the naturally occurring histidine-rich human histatins, have anti-fungal and anti-bacterial activity and are useful in compositions for the treatment of oral, vaginal, urethral, mucosal, respiratory, skin, ear and ophthalmic fungal or bacterial infections. It is particularly effective against local and systemic Candida albicans infection, against oral bacterial diseases such acaries and perphyromonas constitution, and against Clostridium histolyticum. The peptide has superior anti-fungal (especially anti-Candidal) and anti-bacterial activity, particularly on a weight basin, compared to the longer, naturally occurring histatins. Peptides containing D-residues are also more resistant to degradation than L-amino acid versions
                                                                                                                                                                                                                                                                                                                                                                                                                                              - useful esp for
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bacterial disease, Streptococcus mutans; Porphyromonas gingivalis; Clostridium histolyticum.
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    12 /note= "At least one amino acid must have a configuration"

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Pred. No. 3.3e+02;
0; Mismatches 1;
                                                                                            Location/Qualifiers
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75.0%;
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                                                          Synthetic.
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Region
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AAW19454
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The present sequence represents a specifically claimed example of an anti-
-fungal and anti-bacterial D-amino acid histatin-based peptide 113, where
at least one amino acid in the peptide is in the D-configuration. The

peptide is preferably modified by an acetyl or carbamyl addition at the N-
-terminus and/or amidation at the C-terminus. The novel D-amino acid-
containing peptide, based on the naturally occurring histidine-rich human

containing peptide, based on the naturally occurring histidine-rich human

containing peptide, based on the naturally occurring histidine-rich human

compositions for the treatment of oral, vaginal, urethral, mucosal,

respiratory, skin, ear and ophthalmic fungal or bacterial infections. It

is particularly effective against local and systemic Candida albicans

composition, against oral bacterial diseases such as caries and

periodontitis (by inhibiting e.g. Streptococcus mutans and Porphyromonas

gingivalis) and against Clostridium histolyticum. The D-amino acid-
containing peptide has superior anti-fungal (especially anti-candidal)

and anti-bacterial activity, particularly on a weight basis, compared to

the natural L- amino acid forms of histatins and histatin-based peptides.

The presence of D-residues also makes the peptides more resistant to

degradation than corresponding L-amino acid versions
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anti-fungal and anti-bacterial D-amino acid histatins and histatin-based peptide(s) - useful esp for treating Candidal infections, periodontitis
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                                                                                              /note= "At least one amino acid must have D-configuration"
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                                                          Location/Qualifiers
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Matches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                              Xu T,
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                 Synthetic
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fungal and anti-bacterial D-amino acid histatin-based peptide 113-F5,
where at least one amino acid in the peptide is in the D-configuration.

The peptide is preferably modified by an acetyl or carbamyl addition at the N-terminus and/or amidation at the C-terminus. The novel D-amino acid containing peptide, based on the naturally occurring histidine-rich human histatins, have anti-fungal and anti-bacterial activity and are useful in compositions for the treatment of oral, vaginal, urethral,

C useful in compositions for the treatment of oral, vaginal, urethral,

mucosal, respiratory, skin, ear and ophthalmic fungal or bacterial

C andida albicans infection, against oral bacterial diseases such as caries and periodontitis (by inhibiting e.g. Streptococcus mutans and Porphyromonas gingivalis) and against clostridium histolyticum. The D-amino acid-containing peptide has superior anti-fungal (especially anti-candidal) and anti-bacterial activity, particularly on a weight basis, compared to the natural L-amino acid forms of histatins and histatin-based peptides. The presence of D-residues also makes the peptides more resistant to degradation than corresponding L-amino acid versions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents a specifically claimed example of an anti
                                                                                                                                                                                                                                                                                                                                                                                                                              l and anti-bacterial D-amino acid histatins and histatin-based - useful esp for treating Candidal infections, periodontitis
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                                                                                                                                                                                                                                                                                                                                                            Friden PM,
                                                                                        1. .12
/note= "At least one amino acid must have D-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 44.2%; Score 26.5; DB 2; Length 12; 75.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anti-fungal and anti-bacterial D-amino acid peptide 113-F4.
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                                                                                                                                                                                                                                                                                                                                                          Roberts FD,
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                                                                      Location/Qualifiers
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                                                                                                                          configuration"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 2; Fig 1; 63pp; English.
                                                                                                                                                                                                                                 96WO-US009962
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Clostridium histolyticum
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                                                                                                                                                                                                                                                                                                     (PERI-) PERIODONTIX INC.
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                                                                                                                                                                                                                                                                                                                       UNIV BOSTON
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                                                                                                                                                                                                                                                                                                                                                                                                                           Anti-fungal and a peptide(s) - usef and caries.
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                                                                                                                                                         WO9640770-A2
                                                                                                                                                                                                                              1996;
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                                  Synthetic
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AAW19460
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The present sequence represents a specifically claimed example of an anti-
fungal and anti-bacterial D-amino acid histatin-based peptide 113-F4,
where at least one amino acid in the peptide is in the D-configuration.

The peptide is preferably modified by an acetyl or carbamyl addition at
the N-terminus and/or amidation at the C-terminus. The novel D-amino acid
containing peptide, based on the naturally occurring histidine-rich
human histatins, have anti-fungal and anti-bacterial activity and are
useful in compositions for the treatment of oral, vaginal, urethral,
mucosal, respiratory, skin, ear and ophthalmic fungal or bacterial
infections. It is particularly effective against local and systemic
candida albicans infection, against oral bacterial diseases such as
caries and periodontitis (by inhibiting e.g. Streptococcus mutans and
Porphyromonas gingivalis) and against Clostridium histolyticum. The D-
amino acid-containing peptide has superior anti-fungal (especially anti-
candidal) and anti-bacterial activity, particularly on a weight basis,
compared to the natural L- amino acid forms of histatins and histatin-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Anti-fungal and anti-bacterial D-amino acid histatins and histatin-based peptide(s) - useful esp for treating Candidal infections, periodontitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 based peptides. The presence of D-residues also makes the peptides more resistant to degradation than corresponding L-amino acid versions
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bacterial disease; Streptococcus mutans; Porphyromonas gingivalis;
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                                                                                                                         /note= "At least one amino acid must have
configuration"
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Pred. No. 3.3e+02;
0; Mismatches 1;
                                                                                        Location/Qualifiers
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75.0%;
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                  Clostridium histolyticum
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                                                                                                                                                                                                                                                                                                                                                                               Xu T,
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                                                                                                                                                                                                                                                                                                                                          UNIV BOSTON
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Best Local Similarity
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                                                                                                                                                                                                                                                                                        07-JUN-1995;
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                                                                                                                                                                                                                  19-DEC-1996,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and caries.
                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                            UYBO-)
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The present sequence represents a specifically claimed example of an anti-fungal and anti-bacterial D-amino acid histatin-based peptide 113-F4.5, where at least one amino acid in the peptide is in the D-configuration. The peptide is preferably modified by an acetyl or carbamyl addition at the N-terminus and/or amidation at the C-terminus. The novel D-amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            containing peptide, based on the naturally occurring histidine-rich human histatins, have anti-fungal and anti-bacterial activity and are useful in compositions for the treatment of oral, vaginal, urethral, mucosal, respiratory, skin, ear and ophthalmic fungal or bacterial infections. It is particularly effective against local and systemic candida albicans infection, against oral bacterial diseases such as caries and periodontitis (by inhibiting e.g. Streptococcus mutans and Porphyromonas gingivalis) and against clostridium histolyticum. The Damino acid-containing peptide has superior anti-fungal (especially anticandidal) and anti-bacterial activity, particularly on a weight basis, compared to the natural L-amino acid forms of histatins and histatinbased peptides. The presence of D-residues also makes the peptides more resistant to degradation than corresponding L-amino acid versions
                                                                                                                                                                                                                                                                                                                                                                                                                                                       l and anti-bacterial D-amino acid histatins and histatin-based - useful esp for treating Candidal infections, periodontitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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respiratory infection; skin infection; ophthalmic infection; bacterial disease; Streptococcus mutans; Porphyromonas gingivalis; Clostridium histolyticum.
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/note= "At least one amino acid must have

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                                                                                                                                                                                                                                                                                                                                                                                       Roberts FD,
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Pred. No. 3.3e+02;
0; Mismatches 1;
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                                                                                                             Location/Qualifiers
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                                                                                                                                                               configuration"
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                                                                                                                                                                                                                                                                                                                                                                                       Oppenheim FG, Xu T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anti-fungal and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 12 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide (\vec{s}) and caries.
                                                                           Synthetic
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                                                                                                                               Region
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fungal and anti-bacterial D-amino acid histatin-based peptide 113-Q3.9,
where at least one amino acid in the peptide is in the D-configuration.
The peptide is preferably modified by an acetyl or carbamyl addition at the N-terminus and/or amidation at the C-terminus. The novel D-amino acid containing peptide, based on the naturally occurring histidine-rich human histatins, have anti-fungal and anti-bacterial activity and are useful in compositions for the treatment of oral, vaginal, urethral, according in the particularly effective against local and systemic candida albicans infection, against oral bacterial diseases such as caries and periodontitis (by inhibiting e.g. Streptococcus mutans and Porphyromonas gingivalis) and against Clostridium histolyticum. The D-amino acid-containing peptide has superior anti-fungal (especially anti-candidal) and anti-bacterial activity, particularly on a weight basis, compared to the natural L-amino acid forms of histatins and histatin-based peptides. The presence of D-residues also makes the peptides more resistant to degradation than corresponding L-amino acid versions
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anti-fungal and anti-bacterial D-amino acid histatins and histatin-based peptide(s) - useful esp for treating Candidal infections, periodontitis
 mucosal infection; ear infection;
ophthalmic infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
vaginal infection; urethral infection; mucosal infection; ear inferespiratory infection; skin infection; ophthalmic infection; bacterial disease; Streptococcus mutans; Porphyromonas gingivalis; Clostridium histolyticum.
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                                                                                                                                    Score 26.5; DB 2; Length 12;
Pred. No. 3.38+02;
0; Mismatches 1; Indels
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                                                                                                                   Location/Qualifiers
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                                                                                                                                                                   configuration'
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75.08;
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                                                                                                                                                                                                                                                                                                                                                                                        Xu T,
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                    19-DEC-1996
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                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                      (PERI-)
                                                                                                                   Key
Region
                                                                                                                                                                                                                                                                                                                                                        (UYBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 32
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  0 × 2 × 5 × 6 × 6 ×
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This invention describes novel substantially pure antibacterial and antifungal peptides (A) containing between 13 and 20 amino acids, where the peptide has the amino acid sequence: R1-R2-R3-R4-R5-R6-R7-R8-R9-R10-CC R11-R12-R13-R14-R15-R16-R17-R18-R19-R20-R21-R22-R23; where: R1 is Asp or absent; R2 and R20 are Ser or absent; R3, R18, R19 and R21 are His or absent; R4 is Ala; R5, R6, R9, R12, R13, are Lys, G10, Arg or another CC basic amino acid; R7, R8 and R15 are His, Phe, Tyr, Leu, another hydrophobic amino acid, or is absent; R10 is Tyr; R16 is G10 or absent; R17 is Lys or absent; R22 is Arg or absent; R23 is G1y or absent; where CC anino acid sequence. (A) can be used to treat fungal or bacterial amino acid sequence. (A) can be used to treat fungal or bacterial candidal infection, as well as bacterial and viral infections, especially veginal, urethral, mucosal, respiratory, skin, ear, oral, or ophthalmic candida albicans, Actinomyces actinomycetemcomitans, Actinomyces caused by candida albicans, Actinomyces actinomycetemcomitans, Actinomyces viscosus, Bacteriodes forsythus, Bacteriodes gracilis, Bacteriodes vesous, Bacteriodes forsythus, Capnocytophaga gingivalis, Campylobacter sputorum, Capnocytophaga gingivalis, Campylobacter sputorum, Pusobacterium aucideatum, Eusobacterium aucideatum, Fusobacterium constant aucideatum, Fusobacterium constant aucideatum, Pusobacterium aucideatum, Pusobacterium constant aucideatum aucideatum, Eusobacterium aucideatum, Eusopaceatum, Eus
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Antifungal; antibacterial; treatment; infection; candidal; ophthalmic; viral vaginal; urethral; mucosal; respiratory; skin; ear; oral.
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Pred. No. 3.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                              Friden PM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antibacterial and antifungal peptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 7; 31pp; English.
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                                                                                                                                                                                                            98WO-US026513.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  44.2%;
                                                                                                                                                                                                                                                                                                                                                                                Roberts FD, Spacciapoli P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                          PERIODONTIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12
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                                                                                                                                                                                                                                                                                                                               UNIV BOSTON
                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-395161/33
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6 GYKR-KFH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GYKNSKFH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 12 AA;
                                                               Unidentified
                                                                                                               WO9931123-A1
                                                                                                                                                                                                            14-DEC-1998;
                                                                                                                                                                                                                                                            18-DEC-1997;
                                                                                                                                                               24-JUN-1999.
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                                                                                                                                                                                                                                                                                                                                   (UYBO-)
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Unidentified
                                                            24-JUN-1999
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                                                                                                                                                                                                                                                                                                  antifungal peptides (A) containing between 13 and 20 amino acids, where the peptide has the amino acid sequence: R1-R12-R3-R4-R5-R6-R7-R8- R9-R10-R11-R12-R13-R14-R15-R16-R17-R18-R19-R20-R21-R22-R23; where: R1 is Asp or absent; R2 and R20 are Ser or absent; R3, R18, R19 and R21 are His or absent; R4 is Ala; R5, R6, R9, R12, R13, are Lys, Gln, Arg or another basic amino acid; R7, R8 and R15 are His, Phe, Tyr, Leu, another conversed or is absent; R10 is Tyr; R16 is Glu or absent; R17 is Lys or absent; R22 is Arg or absent; R2; R6, R12, and R13 of the amino acid sequence. (A) can be used to treat fungal or bacterial candidal infection, as well as bacterial and viral infections, especially vaginal, urethral, mucosal, respiratory, skin, ear, oral, or ophthalmic infections. The infections that can be treated include those caused by candida albicans, Actinomyces actinomycetemcomitans, Actinomyces viscosus, Bacteriodes gracilis, Bacteriodes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ij
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ureolyticus, Campylobacter concisus, Campylobacter rectus, Campylobacter showae, Campylobacter sputorum, Capnocytophaga gingivalis, Capnocytophaga ochracea, Capnocytophaga sputigena, Clostridium histolyticum, Bikenella
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               corrodens, Bubacterium nodatum, Fusobacterium nucleatum, Fusobacterium peridonticum, Peptostreptococcus nicros, Porphyromonas endodontalis, Porphyromonas gingivalis, Prevotella intermedia, Prevotella nigrescens, Propionibacterium acnes, Pseudomonas aeruginosa, and those caused by various Staphylococcal and Streptococcal species. AAY25140-Y25149 are peptides used in the description of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antifungal; antibacterial; treatment; infection; candidal; ophthalmic; viral vaginal; urethral; mucosal; respiratory; skin; ear; oral.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                          This invention describes novel substantially pure antibacterial and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1;
viral vaginal; urethral; mucosal; respiratory; skin; ear; oral.
                                                                                                                                                                                        Xu T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 26.5; DB 2; Length 12; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
                                                                                                                                                                                        Oppenheim FG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                        Friden PM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antibacterial and antifungal peptide 1.
                                                                                                                                                                                                                                         Antibacterial and antifungal peptides.
                                                                                                                                                                                                                                                                  Disclosure; Page 7; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY25140 standard; peptide; 12 AA
                                                                                                  98WO-US026513.
                                                                                                                          97US-00993235.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 44.2%;
                                                                                                                                                                                        Spacciapoli P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-AUG-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6; Conservative
                                                                                                                                                  (PERI-) PERIODONTIX INC. (UYBO-) UNIV BOSTON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6 GYKR-KFH 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GYKNSKFH 8
                                                                                                                                                                                                                WPI; 1999-395161/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 12 AA;
                        Unidentified
                                                                                                   14~DEC-1998;
                                                                                                                           18-DEC-1997;
                                                                                                                                                                                        Roberts FD,
                                                                          24-JUN-1999.
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This invention describes novel substantially pure antibacterial and antifungal peptides (A) containing between 13 and 20 amino acids, where the peptide has the amino acid sequence: R1-R2-R3-R4-R5-R6-R7-R8- R9-R10-R11-R12-R13-R14-R15-R16-R17-R18-R19-R20-R21-R22-R23; where: R1 is Asp or absent; R2 and R20 are Ser or absent; R3, R18, R19 and R21 are His or absent; R4 is Ala; R5, R6, R9, R12, R13, are Lys, Gln, Arg or another basic amino acid; R7, R8 and R15 are His, Phe, Tyr, Leu, another hydrophobic amino acid, or is absent; R10 is Tyr; R16 is Glu or absent; R17 is Lys or absent; R22 is Arg or absent; R23 is Gly or absent; where Gln cannot simultaneously occupy positions R5, R6, R12, and R13 of the amino acid sequence. (A) can be used to treat fungal or bacterial infection in a mammal. The peptides are particularly used to treat candidal infection, as well as bacterial and viral infections, especially vaginal, urethral, mucosal, respiratory, skin, ear, oral, or ophthalmic infections. The infections that can be treated include those caused by infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Candida albicans, Actinomyces actinomycetemcomitans, Actinomyces viscosus, Bacteriodes forsythus, Bacteriodes gracilis, Bacteriodes ureolyticus, Campylobacter concisus, Campylobacter rectus, Campylobacter showae, Campylobacter sputorum, Capnocytophaga gingivalis, Capnocytophaga ochracea, Capnocytophaga sputigena, Clostridium histolyticum, Eikenella corrodens, Eubacterium nodatum, Fusobacterium nucleatum, Fusobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    peridonticum, Peptostreptococcus nicros, Porphyromonas endodontalis, Porphyromonas gingivalis, Prevotella intermedia, Prevotella nigrescens, Propionibacterium acnes, Pseudomonas aeruginosa, and those caused by various Staphylococcal and Streptococcal species. AAY25140-Y25149 are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Histatin-related peptide; cystic fibrosis; Pseudomonas aeruginosa; antibiotic resistance; pulmonary infection; antibacterial; antibiotic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ij
                                                                                                                                                                                                                                                                                                                                                                                                                           Xu T;
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                                                                                                                                                                                                                                                                                                                                                                                                                           Friden PM, Oppenheim FG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  peptides used in the description of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 26.5; DB 2;
Pred. No. 3.3e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Histatin-related peptide, P-113-R3.9K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antibacterial and antifungal peptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 7; 31pp; English.
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                                                                                                                                                    98WO-US026513.
                                                                                                                                                                                                                                    97US-00993235
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ilarity 75.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                               Roberts FD, Spacciapoli P,
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                                                                                                                                                                                                                                                                                                             (PERI-) PERIODONTIX INC (UYBO-) UNIV BOSTON.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-395161/33
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WO9931123-A1
                                                                                                                                                           14-DEC-1998;
                                                                                                                                                                                                                                    18-DEC-1997;
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Treating cystic fibrosis and combating Pseudomonas and other pulmonary infections in a mammal comprises administering histatin or a fragment of
                              /note= "C-terminal amide"
                Location/Qualifiers
12
                                                                                                                                                   Example 6; Page 13; 27pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                AAB11976 standard; peptide; 12 AA.
                                                                                                     Rothstein DM,
                                                                 07-JAN-2000; 2000WO-US000480
                                                                             99US-00226666
                                                                                        (PERI-) PERIODONTIX INC
                                                                                                                WPI; 2000-465852/40.
                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                       1 GYKNSKFH
                                                                                                                                                                                                                                                                                                                                    Sequence 12 AA;
                                         WO200040204-A2
                  Key
Modified-site
                                                                            08-JAN-1999;
                                                     13-JUL-2000
Homo sapie
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                             AAB11976;
                                                                                                                                                                                                                                                                                                                                                Query Match
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 36
AAB11976
g
                                                                                                                                                                                                                                                                                                                                                                                                                  HXXXH
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The invention relates to treating cystic fibrosis via the administration of a histatin, or a histatin, related peptide. The peptide that is administered contains between 8 and 20 amino acids and has the sequence: X1-X2-X3-X4-X5-X6-X7-X8-X9-X10-X11-X12-X13-X14-X15-X16-X17-X18-X19-X20-X21-X22-X23 where X1 is Asp or absent; X2 is Ser or absent; X3 is His or absent; X4 is Ala, His, Leu, or absent; X5 is Lys, another basic amino acid; X6 is Arg, Gln, Lys, or another basic amino acid; X6 is Arg, Gln, Lys, or another basic amino acid; X6 is Arg, Gln, Lys, or another basic amino acid; X10 is Tyr; X11 is Lys, CT, y15, Ly, Ly, Arg, Ser, or a basic amino acid; X10 is Tyr; X11 is Lys, CT, or another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X13 is Lys, Gln, Arg, Orn, another basic amino acid; X14 is Phe or absent; X15 is His or absent; X16 is Glu or absent; X17 or absent; X21 is His or absent; X22 is Arg or absent; A22 is A23 is Gly or absent. Optionally, the peptide may contain at least one modification and a C-terminal amide addition, an N-terminal carbamyl addition and a C-terminal amide addition, Histatin, histatin fragments armining and histatin-related peptides have activity against Promise Histations and other pulmonary infections in cystic fibrosis patients. Such infections are often resistant to the antibiotics normally used to treat the exemplifications of the invention. With the 7.10 (AABI1988), all these peptides showed aeruginosa. They may therefore be used to combat Pseudomonas infections and other pulmonary infections in cystic fibrosis nationts. pulmonary infections. Sequences AAB11985-B11993 represent relation related peptides used in the exemplifications of the invention. Wit exception of P-113-Q2.3.9.10 (AAB11988), all these peptides showed activity against Pseudomonas aeruginosa

1; 44.2%; Score 26.5; DB 3; Length 12; 75.0%; Pred. No. 3.3e+02; live 0; Mismatches 1; Indels 6; Conservative

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Gaps

6 GYK-KKFH 12

24-NOV-2000 (first entry)

The invention relates to treating cystic fibrosis via the administration of a histatin, a fragment of histatin, or a histatin-related peptide. The peptide that is administered contains between 8 and 20 amino acids and has the sequence: X1-X2-X3-X4-X5-X6-X7-X8-X9-X10-X11-X12-X13-X14-X15-X16-X17-X18-X19-X20-X21-X22-X23 where X1 is Asp or absent; X2 is Ser or absent; X3 is His or absent: X4 is Ala, His, Leu or absent; X5 is Lys, C1n, Arg, Orn, or another basic amino acid; X6 is Arg, G1n, Lys, or another basic amino acid; X7 is His, Phe, Tyr Leu, or another hydrophobic amino acid; X8 is His, Phe, Tyr Leu, or another hydrophobic amino acid; X8 is His, Phe, Tyr Leu, or another hydrophobic amino acid; X10 is Lys, G1n, Arg, C1n, or another basic amino acid; X13 is Lys, G1n, Arg, C1n, or absent; X14 is Phe or absent X15 is His, Phe, Tyr, C1cu, another hydrophobic amino acid, or absent X16 is G10 or absent; X10 is Lys or absent; X10 is His or absent; X20 is Ser or absent; X21 is His or absent; X20 is Arg or absent; X21 is His or absent; X21 is His or absent; X22 is Arg or absent; Andition and a C-terminal acid addition, an N-terminal carbamyl addition, and a C-terminal amide addition, an N-terminal carbamyl addition, and a C-terminal amide addition, Histatin, histatin-related peptides have activity against Pseudomonas and other pulmonary infections in cystic fibrosis patients. Such cuffections are often resistant to the antibiotics normally used to treat or unlmonary infections. Semiances AARI1974-R11844 represent specifically. Treating cystic fibrosis and combating Pseudomonas and other pulmonary infections in a mammal comprises administering histatin or a fragment of pulmonary infections. Sequences AAB11976-B11984 represent specifically claimed histatin-related peptides for use in the method of the invention. The present sequence corresponds to residues 4-15 of naturally-occurring human histatin 3 and histatin 5 (AAB11974, AAB11975) Histatin-related peptide; cystic fibrosis; Pseudomonas aeruginosa; antibiotic resistance; pulmonary infection; antibacterial; antibiotic. Friden PM; Histatin-related peptide, P-113. Claim 4; Page 16; 27pp; English. Rothstein DM, 07-JAN-2000; 2000WO-US000480. 99US-00226666. (PERI-) PERIODONTIX INC. WPI; 2000-465852/40. Spacciapoli P, WO200040204-A2. Sequence 12 AA; Homo sapiens 08-JAN-1999; 13-JUL-2000.

Friden PM;

Gaps 1; Length 12; 1; Indels Score 26.5; DB 3; Pred. No. 3.3e+02; 0; Mismatches 1; 44.2%; 75.0%; 6; Conservative Query Match Best Local Similarity Matches 6; Conserv

1;

GYKR-KFH 12 1 GYKNSKFH 8 셤

AAB11985 standard; peptide; 12 AA. AAB11985; RESULT 37 AAB11985 AXXX AAB11987 standard; peptide; 12 AA.

'note= "N-terminal acetyl" /note= "C-terminal amide"

Location/Qualifiers

RESULT 38 AAB11987

(first entry)

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Treating cystic fibrosis and combating Pseudomonas and other pulmonary infections in a mammal comprises administering histatin or a fragment of
                         Histatin-related peptide; cystic fibrosis; Pseudomonas aeruginosa; antibiotic resistance; pulmonary infection; antibacterial; antibiotic.
             Histatin-related peptide, P-113-NA
                                                                                                                                                                                                                            Example 4; Page 12; 27pp; English.
                                                                                                                                  07-JAN-2000; 2000WO-US000480.
                                                                                                                                                           (PERI-) PERIODONTIX INC.
                                                                                                                                                                                     WPI; 2000-465852/40.
                                                                                                                                                                        Spacciapoli P,
                                                                                                        WO200040204-A2
                                                                       Modified-site
                                                                                    Modified-site
                                                                                                                                               08-JAN-1999;
24-NOV-2000
                                              Homo sapiens
                                                                                                                     13-JUL-2000
                                                     Synthetic.
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Friden PM;

Rothstein DM,

99US-00226666

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activity against Pseudomonas aeruginosa
                                                        Seguence 12 AA;
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ij
                       Gaps
                       ۲,
Length 12;
                      1; Indels
Score 26.5; DB 3;
Pred. No. 3.3e+02;
0; Mismatches 1;
   44.28;
75.08;
                        6; Conservative
   Query Match
Best Local Similarity
                                            1 GYKNSKFH
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Matches

6 GYKR-KFH 12

Length 12;

DB 3;

44.2%; Score 26.5;

activity against Pseudomonas aeruginosa

Sequence 12 AA;

Query Match

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c of a histatin, a tragment of histatin, or a histatin-related peptide. The peptide that is administered contains between 8 and 20 amino acids and has the sequence: X1-X2-X3-X4-X5-X6-X7-X8-X9-X10-X11-X12-X13-X14-X15-X16-X17-X18-X19-X20-X21-X22-X23 where X1 is Asp or absent; X2 is Ser or absent; X3 is His or absent; X4 is Ala, His, Leu or absent; X5 is Lys, CC another basic amino acid; X6 is Arg, Gln, Lys, or another basic amino acid; X6 is Arg, Gln, Lys, Arg, Ser, or a basic amino acid; X10 is Tyr; X11 is Lys, CC amino acid, or another hydrophobic amino acid; X13 is Lys, Gln, Arg, Orn, or another hydrophobic amino acid; X13 is Lys, Gln, Arg, Orn, another hydrophobic amino acid; X13 is Lys, Gln, Arg, Orn, another hydrophobic amino acid, or absent; X14 is Phe or absent; X16 is His or absent; X17, Leu, another hydrophobic amino acid, or absent; X16 is Glu or absent; X20 is Ser or absent; X21 is His or absent; X22 is Arg or absent; and X23 is Gly or absent. Optionally, the peptide may contain at least one modification acid, a build and a C-terminal acyl addition, an N-terminal arglachment and a C-terminal acyl addition. Histatin fragments
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and histatin-related peptides have activity against Pseudomonas and histatin-related peptides have activity against Pseudomonas infections and other pulmonary infections in cystic fibrosis patients. Such infections are often resistant to the antibiotics normally used to treat pulmonary infections. Sequences AAB11985-B11993 represent histatin-related peptides used in the exemplifications of the invention. With the exception of P-113-Q2.3.9.10 (AAB11988), all these peptides showed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Treating cystic fibrosis and combating Pseudomonas and other pulmonary infections in a mammal comprises administering histatin or a fragment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to treating cystic fibrosis via the administration
                                                                                                                                      Histatin-related peptide; cystic fibrosis; Pseudomonas aeruginosa; antibiotic resistance; pulmonary infection; antibacterial; antibiotic.
                                                                                                                                                                                                                                                                                                    /note= "C-terminal amide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Friden PM;
                                                                                                  Histatin-related peptide, P-113-Q3.9.
                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 5; Page 12; 27pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rothstein DM,
                                                                                                                                                                                                                                                                                                                                                                                                                       07-JAN-2000; 2000WO-US000480.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-00226666
                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PERI-) PERIODONTIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-465852/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Spacciapoli P,
                                                                                                                                                                                                                                                                                                                                            WO200040204-A2
                                                                                                                                                                                                                                                                               Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-JAN-1999;
                                                                                                                                                                                                    Homo sapiens.
                                                          24-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                   13-JJL-2000.
                                                                                                                                                                                                                         Synthetic
                   AAB11987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treating
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ή;

Gaps

; ;

1; Indels

Score 26.5; DB 3; Pred. No. 3.3e+02; 0; Mismatches 1;

44.2%;

Local Similarity 75.0 nes 6; Conservative

Matches

Sequence 12 AA;

1;

Query Match

1 GYKNSKFH 8

ð

Length 12;

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The invention relates to treating cystic fibrosis via the administration of a histatin, a fragment of histatin, or a histatin-related peptide. The peptide that is administered contains between 8 and 20 amino acids and 20 mas the sequence: X1-X2-X3-X4-X5-X6-X7-X8-X9-X1-X11-X12-X13-X14-X15-X16-X17-X18-X19-X20-X21-X22-X23 where X1 is Asp or absent; X2 is Ser or absent; X3 is H1s or absent; X4 is Ala, H1s, Leu or absent; X3 is Lys, Cl. or another basic amino acid; X6 is Asy, Gll, Lev, or another basic amino acid; X7 is H1s, Phe, Tyr Leu, or another hydrophobic amino acid; X9 is Gly, Lys, Arg, Ser, or a basic amino acid; X10 is Tyr; X11 is Lys, Cl. or, or another hydrophobic amino acid; X1 is Las, Cl. or another basic amino acid; X1 is Lys, Gln, Arg, Orn, another hydrophobic amino acid; X1 is Lys, Gln, Arg, Orn, another basic amino acid, X1 is Phe or absent X15 is H1s, Phe, Tyr, Leu, or another basic amino acid, X1 is Phe or absent; X10 is Lys, Cl. or absent; X11 is H1s or absent; X14 is H1s or absent; X15 is Gly or absent; X21 is H1s or absent; X19 is H1s or absent; X10 is Ser or absent; X21 is H1s or absent; X19 is H1s or absent; X21 is H1s or absent; X19 is H1s or absent; X19 is H1s or absent; X20 is Ser or absent; X21 is H1s or absent; X19 is H1s or absent; X19 is H1s or absent; X20 is Ser or absent; X21 is H1s or absent; X19 is H1s or absent; X20 is Ser or absent; X21 is H1s or absent;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating cystic fibrosis and combating Pseudomonas and other pulmonary infections in a mammal comprises administering histatin or a fragment of
                                                                                                                                                                                                                                                                                                                                                                                                             Histatin-related peptide; cystic fibrosis; Pseudomonas aeruginosa; antibiotic resistance; pulmonary infection; antibacterial; antibiotic.
                                              Gaps
                                                .,
                                          Indels
. 3.3e+02;
1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "All D-form residues"
                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                        Histatin-related peptide, P-113 (D-form).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Friden PM;
                     Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                    AAB11977 standard; peptide; 12 AA.
                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 5; Page 16; 27pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Spacciapoli P, Rothstein DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JAN-2000; 2000WO-US000480.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-00226666
                75.0%;
                                                                                                                                                                                                                                                                                                                           24-NOV-2000 (first entry)
                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PERI-) PERIODONTIX INC
                                                                                                                          GYK-QKFH 12
                                                                                 1 GYKNSKFH 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-465852/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
Misc-difference 1.
                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200040204-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       )8-JAN-1999;
                                          ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                  AAB11977;
                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to a novel method for peptide synthesis. Specifically, it refers to the synthesis of the protegrin IB-367 peptide by a hybrid process, which comprises carrying out both solid phase synthesis as well as liquid phase synthesis methods. The present invention describes synthesising specific side chain protected peptide fragment intermediates using non-protected arginine reagents, and deprotecting the side chain protected peptide. The protegrin IB-367 peptide is active in vitro and in vivo against microflora, such that it is described as having antibacterial properties and can be used for treating oral mucositis. Furthermore it is useful for coupling to therapeutic agents, reporter molecules, diagnostic agents and antibody molecules. This peptide sequence is the histatin peptide (SeqID 22), synthesised by the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthesizing protegrin IB-367 peptide used for treating infectious diseases, by solid phase synthesis of specific side chain protected peptide, using side chain non-protected arginine reagents and deprotecting side chain protected peptide.
                                                                                                                                                                                                                             peptide synthesis; protegrin IB-367; hybrid process; microflora; antibacterial; oral mucositis; histatin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      44.2%; Score 26.5; DB 7; Length 12; 75.0%; Pred. No. 3.3e+02; iive 0; Mismatches 1; Indels
                                                                                                                                                                                             Biologically active histatin peptide (SeqID 22).
                                                                                        ADB81288 standard; peptide; 12 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 8; 70pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           (INTR-) INTRABIOTICS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Merutka G, Zhu Z, Almeda SA;
                                                                                                                                                                                                                                                                                                                                                                                         22-JAN-2003; 2003WO-US001998.
                                                                                                                                                                                                                                                                                                                                                                                                                         22-JAN-2002; 2002US-0350449P.
                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GYKR-KFH 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-663393/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GYKNSKFH 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                  WO2003062266-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 12 AA;
                                                                                                                                                           04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                     31-JUL-2003
                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                         ADB81288;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                        ADB81288
                                                                                                         g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ð
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Candida albicans; peridontitis; caries; tooth decay; oral infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-SEP-1997
                                                                                                                                                                                                  07-JUN-1996;
                                                                                                                                                        WO9640768-A2
                                                                                                                                                                                                                       07-JUN-1995;
                                                                             Homo sapiens
                                                                                                                                                                             19-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW19452;
                                                                                                              Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 43
AAW19452
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                   Key
g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1;
                                                                                                                                                                                                                                                                                                                                                                                                                        The sequences given in AAR89883-92 are peptide fragments derived from the histidine-rich salivary protein, human and macaque histatin 1 and histatin 3. This sequence represents residues 9-22 of human histatin 3. These peptides are used to treat or prevent infection by Candida albicans in human or veterinary medicine. These peptides are more active than native histatins. These peptides were tested in three assays designed to measure separately their effectiveness in killing of blastoconidia, in killing of germinated cells, and in inhibiting the germination of C. albicans. For these assays, human histatin 3 was isolated from human parotid salivary secretions, and macaque histatin 1 was chemically synthesised. Compositions containing these peptides may also be used in the treatment of bacterial infections, e.g. Streptococcus mutans or P. gingivalis, or viral infections, e.g. Herpes simplex or HIV-1. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                         Treatment of fungal infection with histatins or derived peptide(s) - mostly new cpds., partic. for treating or preventing infection by Candida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                       Histidine-rich salivary protein; HRP; human; macaque; histatin 1; histatin 3; Candida albicans; blastoconidia; inhibition; C. albicans; parotid salivary secretion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .;
;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 26.5; DB 2; Length 14;
Pred. No. 3.9e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anti-fungal and anti-bacterial histatin-based histatin 104.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW19441 standard; peptide; 14 AA
                         AAR89887 standard; peptide; 14 AA.
                                                                                                      Human histatin 3 fragment, 3P4.
                                                                                                                                                                                                                                                                                                                                                                                                       Claim 3; Fig 1; 19pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    44.2%;
                                                                                                                                                                                                                                    94US-00287717.
                                                                                                                                                                                                                                                           91US-00786571.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                    (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6; Conservative
                                                                                                                                                                                                                                                                                           (UYBO-) UNIV BOSTON.
                                                                                                                                                                                                                                                                                                                  Oppenheim FG;
                                                                                                                                                                                                                                                                                                                                      WPI; 1996-097084/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GYKNSKFH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14 AA;
                                                                                                                                                                                                                                                           01-NOV-1991;
28-OCT-1993;
                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                      09-AUG-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-SEP-1997
                                                                                                                                                                                          US5486503-A.
                                                                    25-MAR-2003
23-JUL-1996
                                                                                                                                                                                                                23-JAN-1996,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW19441;
                                                                                                                                                                                                                                                                                                                                                                                   albicans.
                                              AAR89887;
                                                                                                                                                                                                                                                                                                                  Xu T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 42
   RESULT 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW19441
                AAR89887
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The present sequence represents histatin 104, found in human paratoid secretion, where at least one amino acid is in the D-configuration. The secretion, where at least one amino acid is in the D-configuration. The peptide preferably is modified by an acetyl or carbamyl addition at the N-terminus and/or amidation at the C-terminus. The peptide, based on the naturally occurring histidine-rich human histatins, have anti-fungal and are useful in compositions for the treatment of oral, vaginal, urethral, mucosal, respiratory, skin, ear and of oral, vaginal or bacterial infections. It is particularly effective against local and systemic Candida albicans infection, against oral bacterial diseases such as caries and periodontitis (by inhibiting e.g. Streptococcus mutans and Porphyromonas gingivalis) and against clostridium histolyticum. The peptide has superior anti-fungal (especially anti-Candidal) and anti-bacterial activity, particularly on a weight basin, compared to the longer, naturally occurring histatins.

Peptides containing D-residues are also more resistant to degradation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 - useful esp for
                          vaginal infection; urethral infection; mucosal infection; ear infection; respiratory infection; skin infection; ophthalmic infection; bacterial disease; Streptococcus mutans; Porphyromonas gingivalis; Clostridium histolyticum; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
Candida albicans; peridontitis; caries; tooth decay; oral infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1;

    14
/note= "At least one amino acid must have a D-configuration"

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Friden PM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Anti-fungal and anti-bacterial histatin-based peptide(s) treating Candidal infections, periodontitis and caries.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Anti-fungal and anti-bacterial D-amino acid peptide 104.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Spacciapoli P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 26.5; DB 2;
Pred. No. 3.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW19452 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Roberts FD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure, Fig 1; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96WO-US009374.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     44.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95US-00481888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          than L-amino acid versions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PERI-) PERIODONTIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Xu T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ω
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1997-052232/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GYKNSKFH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oppenheim FG,
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Human histatin 3 based peptide 3P4
                                                                                                                                                                                                                                                                                                                    AAW31675 standard; peptide; 14 AA.
                                                                                                                                                   Claim 2; Fig 1; 63pp; English.
                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                         6; Conservative
                                                                                               (PERI-) PERIODONTIX INC
                                                                                                   (UYBO-) UNIV BOSTON
                                                                                                                      WPI; 1997-052234/05
                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                   GYKNSKFH
                                                                                                                                                                                                                                                                                            GYKR-KFH
                                                                                                                                                                                                                                                      Sequence 14 AA;
                                                                                                             Oppenheim FG,
                                                                            07-JUN-1996;
                                                                                     07-JUN-1995;
                                                        WO9640770-A2
                                                                  19-DEC-1996
                                                                                                                                                                                                                                                                                                                                       27-MAR-1998
                                                                                                                                     peptide(s) and caries.
                       Synthetic
                                                                                                                                                                                                                                                                                                                              AAW31675;
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                                     Region
                                                                                                                                                                                                                                                                                                           RESULT 44
                                                                                                                                                                                                                                                                                                               AAW3167
XEXEXEX
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44.2%; 75.0%;

95US-00485273

Xu I,

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This is a human histatin 3 based peptide 3P4. Histatin is a histidine rich salivary protein. This peptide and other peptides based on human histatin 3 and macaque histatins can be synthesised for treatment of fungal, in particular, candidal infections. Expression vectors encoding these peptides can be used in antifungal treatment methods. They can be used to treat candidiasis, fungal infections of the oral cavity, vagina, urethra and skin and systemic fungal infections
              Histatin 3; macaque; human; treatment; fungal infection; candidiasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treating fungal infections - by administering histatin peptide(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human leucocyte antigen; HLA-DQ4; combinatorial library; autoimmune disease; chronic articular rheumatism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 26.5; DB 2;
Pred. No. 3.9e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human leucocyte antigen DQ4 binding peptide #490.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Col 11-12; 19pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW49599 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                   91US-00786571.
93US-00145030.
94US-00287717.
                                                                                                                                                                                                                95US-00441914.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          44.28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94JP-00292657
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6; Conservative
                                histidine rich protein
                                                                                                                                                                                                                                                                                                                                                                         Oppenheim FG, Xu T;
                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-041326/04.
                                                                                                                                                                                                                                                                                                                                 UYBO-) UNIV BOSTON
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GYKNSKFH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GYKR-KFH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 14 AA;
                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUN-1998
                                                                                                                                                                                                              16-MAY-1995;
                                                                                                                                                                                                                                                   11-NOV-10
                                                                                                                                                                                                                                                                                             09-AUG-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP08151396-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-NOV-1994;
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                                                                                                                                 US5696078-A
                                                                                                                                                                        09-DEC-1997
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                                                                                                                                                                                                                                                                        28-OCT-1993
                                                                       Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW49599
              8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents a specifically claimed example of an anticulary and anti-bacterial D-amino acid histatin-based peptide 104, where at least one amino acid in the peptide is in the D-configuration. The peptide is preferably modified by an acetyl or carbamyl addition at the N-terminus and/or amidation at the C-terminus. The novel D-amino acid-containing peptide, based on the naturally occurring histidine-rich human histatins, have anti-fungal and anti-bacterial activity and are useful in compositions for the treatment of oral, vaginal, urethral, mucosal, respiratory, skin, ear and ophthalmic fungal or bacterial infections. It is particularly effective against local and systemic Candida albicans infection, against oral bacterial diseases such as caries and periodontitis (by inhibiting e.g. Streptococcus mutans and Porphyromonas gingivalis) and against Clostridium histolyticum. The D-amino acide containing peptide has superior anti-fungal (especially anti-candidal) and anti-bacterial activity, particularly on a weight basis, compared to the natural L-amino acid forms of histatins and histatin-based peptides. The presence of D-residues also makes the peptides more resistant to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Η;
vaginal infection; urethral infection; mucosal infection; ear infection; respiratory infection; skin infection; ophthalmic infection; bacterial disease; Streptococcus mutans; Porphyromonas gingivalis; Clostridium histolyticum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anti-fungal and anti-bacterial D-amino acid histatins and histatin-based peptide(s) - useful esp for treating Candidal infections, periodontitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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    14
/note= "At least one amino acid must have D-configuration"

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Roberts FD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Spacciapoli P,
                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                            96WO-US009962.
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Gaps

1;

HLA-binding oligopeptide and an immuno:regulator contg it - used in the

WPI; 1996-329479/33

treatment of auto:immune disease.

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0;
                                                                This peptide is an example of a peptide which binds to a human leucocyte antigen HLA-DQ4 molecule. The peptide was isolated from a phagemid combinatorial library comprising the sequence AAV05953, by screening with an HLA-DQ4 molecule. The peptide is used for the treatment of autoimmune disease, or especially for treatment of viral diseases
                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                        .,
                                                                                                                                                                                                                  Query Match 43.3%; Score 26; DB 2; Length 9; Best Local Similarity 44.4%; Pred. No. 1.4e+06; Matches 4; Conservative 1; Mismatches 4; Indels
                                     Claim 4; Page 53; 61pp; Japanese.
                                                                                                                                                                                                                                                                                            1 GYKNSKFHR 9
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1 GYKRDSYER 9
                                                                                                                                                                                   Sequence 9 AA;
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Search completed: August 30, 2004, 10:49:29 Job time : 17.3885 secs

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Q865c9 sus scrofa
Q9au99 aethionema
Q9h326 homo sapien
Q9ps56 carassius a
P87031 saccharomyc
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Q9t4k5 bryopsis sp
Q9t4k4 bryopsis sp
Q9t4k4 bryopsis sp
Q9t4k6 bryopsis sp
Q9t4k3 bryopsis sp
Q9t4k3 bryopsis sp
Q9tum9 homo sapien
Q9lcsl bacillus su
Q9lcsl aedes aegyp
Q9r3u3 chlamydia t
                                                                           August 30, 2004, 10:40:34; Search time 200.27 Seconds (without alignments) 327.696 Million cell updates/sec
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                                                                                                                                 US-09-720-469A-44
1083
1 MKVLLAAALIAGSVFFLLLP......VIIADCGKIEVEKPFAIAKE 208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Description
                                                                                                                                                                                                                                                              3347
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                 1017041 seqs, 315518202 residues
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Maximum Match 100%
Listing first 45 summaries
                                                   OM protein - protein search, using sw model
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Q9AU99
Q9H326
Q9P326
Q9P031
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Gapop 10.0 , Gapext 0.5
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Match Length DB
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Q26100 pratylenchu	Q9nlv6 equus cabal		Oggve9 mus sp. pro	O9n2b9 gorilla gor	09n2b8 pondo pydma	O9n2c0 pan troglod	093vc8 nicotiana t	08qsb9 lolium pere	Q80xv4 rattus sp.	Q64313 rattus norv	Д	P82848 rana pipien		O8jdm7 human immun	OBidmo human immun	O9ths3 bryopsis sp	Q84f20 bacillus su	Q85q96 pyrrhobryum	Q8sac2 amblysteqiu	047606 escherichia	Q9c057 homo sapien	Q9bqq1 homo sapien	Q9xnr6 pylaiella 1	Q8h6e6_hordeum vul	Q80iq7 choristoneu				ı
Q26100	9AIN6O	Q52636	Q9QVE9	Q9N2B9	Q9N2B8	Q9N2C0	Q93YC8	Q8GSB9	Q80XV4	Q64313	072526	P82848	овломз	Q8JDM7	QBJDM0	Q9THS3	Q84F20	982380	Q8SAC2	047606	Q9C057	Q9BQ01	Q9XNR6	Q8H6E6	Q80IG7	O7ZZN7	Q98YM1	Q99PB8	
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2.5	2.5	2.4	2.4	2.4		2.4	2.4	2.4	2.4	2.4	2.4	2.4	2.4	2.4	2.4	•	•	•	•	•	2.3	•		٠	•	•	•	•	
27	27	26.5			26		26				26		26	56	26	25.5	25	25	25	25	25	25	25	25	25	25	25	25	
17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32			32		37	38	39		41	42	43	44	45	

ALIGNMENTS

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01-MAY-2000
01-DEC-2001
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P87031
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Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
Cyprinidae; Carassius.
ICSE_TaxID=7957;
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;
eurosids II; Brassicales; Brassicaceae; Aethionema.
NCBI_TaxID=72657;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Kim Y.-O., Koh H.-J., Jo S.-H., Son M.-K., Huh T.-L.;

"Structural and functional characterization of the human NAD+-
                                                                                                            SEQUENCE FROM N.A.

Koch M., Kroymann J., Haubold B., Weisshaar B., Mitchell-Olds
"Phylogenetic analysis of promoter sequences from cruciferous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 dependent isocitrate dehydrogenase alpha subunit promoter."; Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AF157515; AAG43379.1; ...
                                                                                                                                                                                                                                                                                            Score 32; DB 10; Length 13;
Pred. No. 2e+04;
2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.9%; Score 31; DB 4; Length 9; 66.7%; Pred. No. 1e+06; tive 1; Mismatches 1; Indels
                                                                                                                                                                           plants.";
Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
NAD+-dependent isocitrate dehydrogenase 3 alpha subunit
                                                                                                                                                                                                                                                       13 AA; 1432 MW; 5C444963C4621AA7 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
NEUROLIN=CELL surface glycoprotein (Fragment).
Carassius auratus (Goldfish).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9 AA; 960 MW; C91CB0437DC7687D CRC64;
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Paschke K.A., Lottspeich F., Stuermer C.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
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60.0%;
                                                                                                                                                                                                                 EMBL; AF249000; AAK31935.1;
NON TER 13 13
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                                                                                                                                                                                                                                                                                                                                                                                                         |||: || :|
2 GPSSLDEIRK 11
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GPAWIS
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Q9PS56
ID Q9PS
AC Q9PS
DT 01-M
DT 01-M
DT 01-M
DE NEUR
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          growing retinal axons in the
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Yoshikawa T., Turner G., Esterling L.E., Sanders A.R.,
Detera-Wadleigh S.D.;
"A novel human myo-inositol monophosphatase gene, IMP.18p, maps to a susceptibility region for bipolar disorder.";
Mol. Psychiatry 2:393-397(1997).
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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"Neurolin, a cell surface glycoprotein on growing retinal axigoldfish visual system, is reexpressed during retinal axonal regeneration.";
J. Cell Biol. 117:863-875(1992).
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Van Dyck L., Skala J., de Wergifosse P., Purnelle B., Talla
Nawrocki A., Del Bino S., Goffeau A.;
Submitted (MAY-1996) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                     Score 31; DB 13; Length 13; Pred. No. 2.4e+04; 4; Mismatches 1; Indels
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                                                                                                                                            33AF8FF8CC535728 CRC64;
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(TrEMBLrel. 12, Last sequence update)
(TrEMBLrel. 22, Last annotation update)
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Last annotation update)
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Pred. No. 3.6e+04;
1; Mismatches 1;
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01-OCT-2002 (TrEMBLrel
ORF YGR126W (Fragment)
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SEQUENCE FROM N.A.
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STRAIN=VARIOUS STRAINS;
Krellwitz E.C., Kowallik K.V., Manos P.S.;
"Phylogenetic Study of Bryopsis (Chlorophyta) from the Western North Atlantic and Caribbean based on Coding and Non-coding sequences of the
                      MEDLINE=20284187;
Yoshikawa T., Padigaru M., Karkera J.D., Sharma M., Berrettini W.H.,
Esterling L.E., Detera-Wadleigh S.D.;
"Genomic structure and novel variants of myo-inositol monophosphatase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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Bryopsidaceae; Bryopsis.
NCBI_TaxID=103786;
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Submitted (JUL-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AF170407; AAD56846.1; -.
EMBL; AF170383; AAD56800.1; -.
EMBL; AF170385; AAD56802.1; -.
EMBL; AF170385; AAD56802.1; -.
EMBL; AF170386; AAD56804.1; -.
EMBL; AF170386; AAD56806.1; -.
EMBL; AF170386; AAD56806.1; -.
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01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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EMBL; AF170390; AAD56812.1; -.
EMBL; AF170391; AAD56814.1; -.
EMBL; AF170392; AAD56816.1; -.
EMBL; AF170393; AAD56818.1; -.
EMBL; AF170394; AAD56820.1; -.
GO; GO:0009507; C:chloroplast; IEA.
                                                                                                                                             Mol. Psychiatry 5:165-171(2000).

EMBL; AF025882; AAD22136.1; -.

EMBL; AF025881; AAD22136.1; JOINED.

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SEQUENCE FROM N.A.

STRAIN=VARIOUS STRAINS;

RY Krellwitz E.C., Kowallik K.V., Manos P.S.;

RT Alantic and Caribbean based on Coding and Non-coding sequences of the Chloroplast psbB Operon.";

RI Atlantic ARIO405; AAD56842.1; -.

REMBL; ARI70395; AAD56822.1; -.

REMBL; ARI70395; AAD56824.1; -.

REMBL; ARI70396; AAD5682.1; -.

REMBL; ARI70396; AAD5682.1; -.

REMBL; ARI70396; AAD5683.1; -.

REMBL; ARI70399; AAD5683.1; -.

REMBL; ARI70400; AAD5683.1; -.

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Krellwitz E.C., Kowallik K.V., Manos P.S.;
"Phylogenetic Study of Bryopsis (Chlorophyta) from the Western North Atlantic and Caribbean based on Coding and Non-coding sequences of the Chloroplast psbB Operon.";
Submitted (JUL-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AF170406; AAD56844.1; -.
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Chloroplast.
Eukaryota; Viridiplantae; Chlorophyta; Ulvophyceae; Caulerpales;
Bryopsidaceae; Bryopsis.
NCBI_TaxID=103785;
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Bryopsidaceae; Bryopsis.
NCBI_TaxID=103785;
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SEQUENCE 13 AA; 1351 MW; 2C316AF02E93A338 CRC64;
                                               (TrEMBLrel. 13, Created)
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(TrEMBLrel. 24, Last annotation update)
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01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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STRAIN='VARIOUS STRAINS;
Krellwitz E.C., Kowallik K.V., Manos P.S.;
Krellwitz E.C., Kowallik K.V., Manos P.S.;
"Phylogenetic Study of Bryopsis (Chlorophyta) from the Western North Atlantic and Caribbean based on Coding and Non-coding sequences of the Chloroplast psbB Operon.";
Submitted (JUL-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AF170415; AAD56848.1; -.
EMBL; AF170409; AAD56850.1; -.
EMBL; AF170410; AAD56852.1; -.
EMBL; AF170410; AAD56852.1; -.
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Krellwitz E.C., Kowallik K.V., Manos P.S.;
"Phylogenetic Study of Bryopsis (Chlorophyta) from the Western North
Atlantic and Caribbean based on Coding and Non-coding sequences of the
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Bryopsidaceae; Bryopsis.
NCBI_TaxID=103784;
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Eukaryota; Viridiplantae; Chlorophyta; Ulvophyceae; Caulerpales;
Bryopsidaceae; Bryopsis.
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       Score 29; DB 8; Length 13;
Pred. No. 3.6e+04;
1; Mismatches 0; Indels
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Submitted (JUL-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AF170382; AAD56796.1; -.
GO; GO:0009507; C:chloroplast; IEA.
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Last annotation update)
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                                                                     5; Conservative
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Best Local Similarity
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       Query Match
Best Local Similarity
Matches 5; Conserv
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Q9THS2
ID Q9THS2
DT 01-MA
DT 01-MA
DT 01-JU
DB PBSH.
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STRAIN=95-05-03, 96-07-03, 95-12-02, 94-19-1R, 97-13-1R, and 98-08-03;

Krellwitz E.C., Kowallik K.V., Manos P.S.;

Krellwitz E.C., Kowallik K.V., Manos P.S.;

"Phylogenetic Study of Bryopsis (Chlorophyta) from the Western North

Atlantic and Caribbean based on Coding and Non-coding sequences of the

Chloroplast psbB Operon.";

Submitted (JUL-1999) to the EMBL/GenBank/DDBJ databases.

EMBL; AF170373; AAD56798.1; -.

BMBL; AF170375; AAD5678.1; -.

RMBL; AF170377; AAD56788.1; -.

RMBL; AF170377; AAD56788.1; -.

RMBL; AF170377; C:chloroplast; IEA.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Chloroplast.
Eukaryota; Viridiplantae; Chlorophyta; Ulvophyceae; Caulerpales;
Bryopsidaceae; Bryopsis.
NCBI_TaxID=103788;
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                                                                                                                                       2C31687A2E93A338 CRC64;
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Last sequence update;
Last annotation update)
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EMBL; AF170412; AAD56856.1; -.
EMBL; AF170414; AAD56860.1; -.
GO; GO:0009507; C:chloroplast; IEA.
Chloroplast.
NON TER
13
SEQUENCE 13 AA; 1366 MW; 2C3168
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01-MAY-2000 (TrEMBLrel. 13,
01-MAY-2000 (TrEMBLrel. 13,
01-OCT-2003 (TrEMBLrel. 25,
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STRAIN=Rockefeller/Red; TISSUE=Salivary gland;
MEDLINE=20099025; PubMed=10620041;
Beerntsen B.T., Champagne D.E., Coleman J.L., Campos Y.A., James A.A.;
"Characterization of the Sialokinin I gene encoding the salivary vasodilator of the yellow fever mosquito, Aedes aegypti.";
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Sialokinin I preproprotein (Fragment).
Aedes aegypti (Yellowfever mosquito).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea; Aedes.
NET TaxID=7159;
Keddache M, Zhang F, Greenberg D.A.;
"A Dinucleotide Repeat Between Exons 2 and 3 of the Human RING3 Gene.";
Submitted (NOV-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF107699; AAD24963.1; -.
InterPro; IPR001487; Bromodomain.
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"Identification of IS210 in Azotobacter vinelandii: a novel, functional insertion element member of the IS5 family.";
Submitted (SEP-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ249381; CAB76429.1; -.
GO; GO:0046821; C:extrachromosomal DNA; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Natch 2.6%; Score 28; DB 2; Length 14; Local Similarity 54.5%; Pred. No. 4.9e+04; es 6; Conservative 1; Mismatches 4; Indels
                                                                                                                                                                                                                   Score 29; DB 4; Length 14;
Pred. No. 4e+04;
3; Mismatches 4; Indels
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NCBI_TaxID=1423;
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Last annotation update)
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14 AA; 1677 MW;
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14 AA; 1623 MW;
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Best Local Similarity 41.7%;
Matches 5; Conservative
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Eukaryota; Metazoa; Nematoda; Chromadorea; Tylenchida; Tylenchina;
Tylenchoidea; Pratylenchidae; Pratylenchinae; Pratylenchus.
NCBI_TaxID=45929;
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STRAIN=Maryland isolate;
Setterquist R.A., Smith G.K., Jones R., Fox G.E.;
"Diagnostic Probes Targeting the Major Sperm Protein Gene may be Useful for the Molecular Identification of Nematodes.";
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Hackstadt T.;
"Purification and N-terminal amino acid sequences of Chlamydia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
27 kDa histone analog (Fragment).
Chlamydia trachomatis.
Bacteria; Chlamydiae; Chlamydiales; Chlamydia.
                                                                                                                                                                                                Score 27; DB 5; Length 11; Pred. No. 4.4e+04; 1; Mismatches 5; Indels
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Insect Mol. Biol. 8:459-467(1999).

EMBL; AF108100; AAD16884.1; -.

GO; GO:0007268; P:synaptic transmission; IEA.

GO; GO:0007217; P:tachykinin signaling pathway; IEA.

InterPro; IPR002040; Tachy Neurokinin.

PROSITE; PS00267; TACHYKININ; 1.

NON TER

1 1 A; 1203 MW; 8BADC77C6B59C33A CRC64;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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(TrEMBLrel. 13, Last sequence update)
(TrEMBLrel. 21, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  trachomatis histone analogs.";
J. Bacteriol. 173:7046-7049(1991).
NON TER 14
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PPMSP-5.
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Matches
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Q9R3U3
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Q9N2B9
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Caetano A.R., Shiue Y.L., Lyons L.A., O'Brien S.J., Laughlin T.F.,
Bowling A.T., Murray J.D.;
"A comparative gene map of the horse (Equus caballus).";
Genome Res. 9:1239-1249(1999).
EMBL; AF134224; AAF63865.1; -.
NON_TER 1 1 1
NON_TER 14 14
SEQUENCE 14 AA; 1456 MW; 5C14536E1EFF1977 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Perissodactyla, Equidae, Equus.
NCBI_TaxID=9796;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Plasmid R124.
Bacteria, Proteobacteria, Gammaproteobacteria, Enterobacteriales,
Enterobacteriaceae, Escherichia.
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Frost L.S., Finlay B.B., Opgenorth A., Paranchych W., Lee J.S.;
"Characterization and sequence analysis of pilin from F-like
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 27; DB 6; Length 14;
Pred. No. 6e+04;
4; Mismatches 2; Indels
  2.5%; Score 27; DB 5; Length 14; 36.4%; Pred. No. 6e+04;
                                                                 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                    01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
Beta-hemoglobin (Fragment).
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Last sequence update)
Last annotation update)
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GO: GO:0046821; C:extrachromosomal DNA; IEA.
                                                                                                                                                                                                                                                                                                                                                   14 AA
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                                                            3; Mismatches
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                                                                 4; Conservative
                                                                                                                            155 DGKHVVFGKVL 165
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                                                                                                                                                                         DDKHTYYMKII 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Equus caballus (Horse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tral protein (Fragment).
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                              Best Local Similarity
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SEQUENCE
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Query Match
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Q52636
ID Q52636
AC Q5263
AC Q5263
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Q9N1V6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE-92174934; PubMed=1541289;
MEDLINE-92174934; PubMed=1541289;
Chauviere M., Martinage A., Debarle M., Sautiere P., Chevaillier P.;
Chauviere daracterization of six intermediate proteins in the processing of mouse protamine P2 precursor.";
Eur. J. Biochem. 204:759-765(1992).

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NON_TER 10 10

SEQUENCE 10 AA; 1029 MW; FCAD2DD8676866D1 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                        Mammalia, Butheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Gorilla.
NCBI_TaxID=9593;
                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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80.0%; Pred. No. 4.7e+04;
iive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.4%; Score 26; DB 6; Length 12; 71.4%; Pred. No. 6e+04; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=gorilla-U1;
Kitano T., Kobayakawa H., Saitou N.;
"Silver Project.";
Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AB041364; BAA94453.1;
GO; GO:0008168; F:methyltransferase activity; IEA.
GO; GO:0016740; F:transferase activity; IEA.
                                                                                                                                                                                                                        01_MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Protamine MP2 intermediate protein PMP2/11 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10 AA; 1029 MW; FCAD2DD8676866D1 CRC64;
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12 AA; 1289 MW; CF0798E2917861A9 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Indolethylamine N-methyltransferase (Fragment).
                                                                                                                                                                             10 AA
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NON TER 12
SEQUENCE 12 AA; 1289 MW; CF
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Conservative
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                                                                                                                                                                             PRELIMINARY;
117 DENFKLKHY 125
                                                 4 DEN-KLKKY 11
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es 5; Conserv
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                         01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Putative coat protein (Fragment).
Nicotiana tabacum (Common tobacco).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids; lamiids; Solanales; Solanaceae; Nicotiana.
NCEL TaxID=4097;
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooideae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "ENOD40 homolog from perennial ryegrass (Lolium perenne)."; Submitted (AUG-2002) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.4%; Score 26; DB 10; Length 12; 100.0%; Pred. No. 6e+04; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.4%; Score 26; DB 10; Length 12; 40.0%; Pred. No. 6e+04; ive 3; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=20027532; PubMed=10557305; Jakowitsch J., Mette M.F., van der Winden J., Matzke M.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "ENOD40 G:; "; submitted (AUG-2002) to the EMBL/GenBank/DDBJ databases. EMBL; AF538350; AAN15133.1; -. EMBL; AF538351; AAN15134.1; -. SEQUENCE 12 AA; 1455 MW; 3EF4B0A918EB1733 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
van der Winden J.;
Submitted (SEP-2001) to the EMBL/GenBank/DDBJ databases.
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12 AA; 1402 MW; 89226B3A0351E321 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      dispersed repetitive DNA in plants.",
Proc. Natl. Acad. Sci. U.S.A. 96:13241-13246(1999).
EMBL; AJ414170; CAC88799.1; -.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule activity; IEA.
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
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Larsen K.;
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Les 4; Conservative
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Pan.
                                                                                                                                                                                                                                                                                                                                                                   rongo pygmaeus (Urangutan).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pongo.
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STRAIN=chimp-220;
Kitano T., Kobayakawa H., Saitou N.;
"Silver Project.";
Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases.
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"Silver Project.";
Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases.
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01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Indolethylamine N-methyltransferase (Fragment).
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GO; GO:0008168; F:methyltransferase activity; IEA.
GO; GO:0016740; F:transferase activity; IEA.
Methyltransferase; Transferase.
NON TER 12 12 12
SEQÜENCE 12 AA; 1289 MW; CF0798E2917861A9 CRC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; AB041365; BAA94454.1; -. GO: GO:0008168; F:methyltransferase activity; IEA GO; GO:0016740; F:transferase activity; IEA.
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NON TER 12 12
SEQÜENCE 12 AA; 1290 MW; CF(
                                                                                                                                                                                                                                                                                                                                                             Pongo pygmaeus (Orangutan)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
NCBI_TaxID=10118;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rattus norvegicus (Rat).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
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Rao S.M., Howells R.D.;
"cis-acting elements in the 5'-untranslated region of rat testis proenkephalin mRNA regulate translation of the precursor protein.";
J. Biol. Chem. 268:22164-22169(1993).
EMBL; S66180; AAP13974.1; -.
SEQUENCE 12 AA; 1436 MW; D33C6D2A83B36776 CRC64;
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MEDLINE=90066445; PubMed=2573832;
Garrett J.E., Collard M.W., Douglass J.O.;
"Translational control of germ cell-expressed mRNA imposed by alternative splicing: opioid peptide gene expression in rat te Mol. Cell. Biol. 9:4381-4389(1989).
EMBL; K02805; AAA60733.1; -.
                                                                                                                                                                                                                                                                                             2.4%; Score 26; DB 11; Length 12; 66.7%; Pred. No. 6e+04; tive 0; Mismatches 2; Indels
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Hypothetical protein.
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                              12 AA.
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                                                                                 Created)
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Q7Z5Z6;
01-OCT-2003 (TrEMBLrel. 25, Created)
                                                         PRT;
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                                                                            (TrEMBLrel. 24, C
TrEMBLrel. 24, I
(TrEMBLrel. 24, I
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Best Local Similarity 66.,
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Best Local Similarity 66.7
Matches 4; Conservative
                                                        PRELIMINARY;
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                                                                                         01-JUN-2003 (TrEMBLrel. 01-JUN-2003 (TrEMBLrel.
MEDAWLEHLH 10
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                                                                                -JUN-2003
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                                                                  Q80XV4;
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Q7Z5Z6
ID Q7Z5Z0
AC Q7Z5Z0
DT 01-OC
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                                                                                                                                                                                                                          Shu J., Li G., He X.; "Construction of cDNA expression library from nasopharyngeal carcinoma tissue and screening of antigenic genes."; Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases. EMBL; AY320404; AAP73806.1; -. SEQUENCE 13 AA; 1429 MW; 1D0C143838017694 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Peptides with antimicrobial activity from four different families isolated from the skins of the North American frogs Rana luteiventris, Rana berlandieri and Rana pipiens.";

Bur. J. Biochem. 267:894-900(2000),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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-!- MASS SPECTROMETRY: MW=1368; METHOD=ELECTROSPRAY.
-!- SIMILARITY: BELONGS TO THE BREVININ/ESCULENTIN/GAEGURIN/RUGOSIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Ranoidea, Ranidae, Rana.
NCBI_TaxID=8404;
                                                                   Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Goraya J., Wang Y., Li Z., O'Flaherty M., Knoop F.C., Platz J.E., Conlon J.M.;
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                                                                                                                                                                                                                                                                                                                                                                                                               Score 26; DB 4; Length 13; Pred. No. 6.6e+04; 1; Mismatches 1; Indels
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3EF3402B9DF92338 CRC64;
25, Last sequence update) 25, Last annotation update)
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(TrEMBLrel. 16, Last sequence update)
(TrEMBLrel. 24, Last annotation update)
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rana pipiens (Northern leopard frog)
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MEDLINE=20117700; Pubmed=10651828;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      71.4%;
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(TrEMBLrel. 22, I
(TrEMBLrel. 25, I
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SEQUENCE FROM N.A.
STRAIN=96-21-02;
Krellwitz E.C., Kowallik K.V., Manos P.S.;
"Phylogenetic Study of Bryopsis (Chlorophyta) from the Western North Atlantic and Caribbean based on Coding and Non-coding sequences of the Chloroplast psbB Operon.";
Submitted (JUL-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AF170381; AAD56794.1; -.
MEDLINE=22032414; PubMed=12036486;
Long E.M., Rainwater S.M., Lavreys L., Mandaliya K., Overbaugh J.;
"HIV Type I Variants Transmitted to Women in Kenya Require the CCRS
Coreceptor for Entry, Regardless of the Genetic Complexity of the
Infecting Virus.";
AIDS Res. Hum. Retroviruses 18:567-576(2002).
EMBL; AF407150; AAM66205.1; -.
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MEDLINE=89034018; PubMed=2460441;
Nakamura K., Itoh Y., Yamane K.;
"Enhanced secretion of beta-lactamase on structural modification of the Bacillus subtilis alpha-amylase signal peptide.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Viridiplantae; Chlorophyta; Ulvophyceae; Caulerpales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 25.5; DB 8; Length 13;
Pred. No. 7.3e+04;
1; Mismatches 0; Indels
                                                                                                                                                                                           2.4%; Score 26; DB 15; Length 14; 83.3%; Pred. No. 7.2e+04; tive 0; Mismatches 1; Indels
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Bacillus subtilis.
Bacteria, Firmicutes; Bacillales; Bacillaceae; Bacillus.
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13 AA; 1379 MW; 2C31687A3B73A338 CRC64;
                                                                                                                                                        14 AA; 1442 MW; 37C58F7BF82D7AA8 CRC64;
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Last annotation update)
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01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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Local Similarity 75.0%;
hes 6; Conservative
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les 5; Conservative
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NCBI_TaxID=103787;
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                                                                                                                                                        SEQÜENCE
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Q84F20;
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Q84F20
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Matches
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                                                                                                SEQUENCE FROM N.A.

MEDLINE=22032414; PubMed=12036486;
Long E.M., Rainwater S.M., Lavreys L., Mandaliya K., Overbaugh J.;
Long E.M., Rainwater S.M., Lavreys L., Mandaliya K., Overbaugh J.;
"HIV Type 1 Variants Transmitted to Women in Kenya Require the CCR5
"HIV Type 1 Variants Transmitted to Women in Kenya Require the CCR5
"Coreceptor for Entry, Regardless of the Genetic Complexity of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE=22032414; PubMed=12036486;
Long E.M., Rainwater S.M., Lavreys L., Mandaliya K., Overbaugh J.;
"HIV Type 1 Variants Transmitted to Women in Kenya Require the CCR5
Coreceptor for Entry, Regardless of the Genetic Complexity of the
Infecting Virus.";
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Pred. No. 7.2e+04;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                           2.4%; Score 26; DB 15; Length 14; 83.3%; Pred. No. 7.2e+04; tive 0; Mismatches 1; Indels
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01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
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01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Truncated tat protein (Fragment).
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
                                          Human immunodeficiency virus 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
VCBI_TaxID=11676;
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NCBI_TaxID=11676,
                                                                                                                                                                                                            Infecting Virus.";
AIDS Res. Hum. Retroviruses 18:567-576(2002)
EMBL; AF407149; AAM66202.1; -.
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         Truncated tat protein (Fragment).
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Best Local Similarity 83.33
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Best Local Similarity 83.3
Matches 5; Conservative
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                                                                          5; Conservative
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Best Local Similarity 71.4
Matches 5; Conservative
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                                                 Best Local Similarity
Matches 5; Conserv
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Best Local Similarity
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SEQUENCE FROM N.A.
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Q47606
ID Q4760
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"Reconciled gene trees, molecular evolution, and species relationships in Amblystegium (Bryopsida).";
Submitted (JAN-2002) to the EMBL/GenBank/DDBJ databases.

EMBL; AF464972; AAL75469.1; -..

NON TER 10 10 10 10 10 SEQUENCE 10 AA; 1143 MW; 9D5040D1B415BAB1 CRC64;
                                                                                                                                                                                 Gaps
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Amblystegium varium.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Bryophyta;
Bryopsida; Bryidae; Hypnales; Amblystegiaceae; Amblystegium.
                                                                                                                                                                               ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chloroplast.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Bryophyta;
Bryopsida; Bryidae; Rhizogoniales; Rhizogoniaceae; Pyrrhobryum.
NCBI_TaxID=129293;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDINE=22569835; PubMed=12683518;
McDaniel S.F., Shaw A.J.;
"Phylogeographic structure and cryptic speciation in the trans-
Antarctic moss Pyrrhobryum mnioides.";
Evolution 57:205-215(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
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                                                                                                                             Query Match 2.3%; Score 25; DB 2; Length 9; Best Local Similarity 71.4%; Pred. No. 1e+06; Matches 5; Conservative 1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                             Created)
Last sequence update)
Last annotation update)
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01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                   9 9 AA; 728 MW; C6B9BDDDDDC7687D CRC64;
                                                                                                                                                                                                                                                                                                                                                                    9 AA.
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                                                                                                                                                                                                                                                                                                                                                                      PRT;
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J. Biochem. 104:265-269(1988).

EMBL; D13786; BAC56113.1; -.

NON TER 1 1:

NON_TER 9 9

SEQUENCE 9 AA; 728 MW; C6B9
                                                                                                                                                                                                                                                                                                                                                                                                          01-JUN-2003 (TrEMBLrel. 24, 01-JUN-2003 (TrEMBLrel. 24, 01-OCT-2003 (TrEMBLrel. 25,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pyrrhobryum mnioides.
                                                                                                                                                                                                                     19 LPGPSAA 25
                                                                                                                                                                                                                                                             1 LAGPAAA 7
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nes 5; Conserv
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ATPB.
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SEQUENCE
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Q85G96
ID Q85G9
AC Q85G9
DT 01-JU
DT 0
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Q8SAC2
ID Q8SAC2
DT Q8SAC3
DT O1-JU
DT O1-JU
DT O1-JU
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OS AMDIY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=91139577; PubMed=1995588;
Tao T., Bourne J.C., Blumenthal R.M.;
"A family of regulatory genes associated with type II restriction-modification systems.";
J. Bacteriol. 173:1367-1375(1991).
EMBL; M63622; AAA24562.1;
NON TER. 11 11.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
Enterobacteriaceae; Escherichia.
NCBI_TaxID=562;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TAXID=9606;
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Submitted (SEP-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AF182950; AAK12833.1; -.
NON TER 11
SEQUENCE 11 AA; 1125 MW; 2644D7FE686761F7 CRC64;
2.3%; Score 25; DB 10; Length 10; 100.0%; Pred. No. 5.8e+04; tive 0; Mismatches 0; Indels
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Pred. No. 6.5e+04;
1; Mismatches 1; Indels
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01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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Last sequence update)
Last annotation update)
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80.0%; Pred. No. 6.5e+04;
ative 1; Mismatches 0;
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SEQUENCE FROM N.A.
TISSUE=Cerebellum;
Weber J.R., Sokol S.Y.;
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NCBL_TaxID=8355;
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                                                                                                         STRAIN=cv. Orthega;
Larsen K.;
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Best Local Similarity
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                                                                                           SEQUENCE FROM N.A.
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                                                                         NCBI_TaxID=4513;
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Best Local 9
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Q801G7;
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"The mitochondrial Pylaiella littoralis nad11 gene contains only the N-terminal FeS-binding domain.";
Gene 235:131-137(1999).
EMBL; AF110139; AAD44051.1; -.
GO; GO:0005739; C:mitochondrion; IEA.
Mitochondrion; Ubiquinone.
                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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45.5%; Pred. No. 7.3e+04;
cive 1; Mismatches 5; Indels
                                                                                                                                                                                                                                      Score 25; DB 4; Length 12;
Pred. No. 7.3e+04;
0; Mismatches 1; Indels
                                                                                                                                                                        Lloyd D.;
Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AL121582; CAC36069.1; -.
NON_TER 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; stramenopiles; Phaeophyceae; Ectocarpales;
Acinetosporaceae; Pylaiella.
NCBL_TaxID=2885;
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12 AA; 1501 MW; 49750746424B5B13 CRC64;
                                                                                                                                                                                                                  SEQÜENCE 12 AA; 1017 MW; 4789B7218E787877 CRC64;
                                        09B001;
01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
BA28009.2.2 (Novel protein (Isoform 3)) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                               01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
NADH:ubiquinone oxidoreductase subunit 3 (Fragment)
                                                                                                                                                                                                                                                                                                                                                             12 AA.
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                              12 AA.
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01-MAR-2003 (TrEMBLrel. 23, Created)
                               PRT;
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Best Local Similarity 83.3%;
Matches 5; Conservative
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                               PRELIMINARY;
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                                                                                                          Homo sapiens (Human)
                                                                                                                                                                                                                                                                                  104 GTGGKS 109
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Matches 5; Conserv
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                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                         NCBI_TaxID=9606;
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                                                                                              DJ631M13.5
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SEQUENCE
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                              09BQQ1
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         RESULT 39
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ID Q8
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                                                                                       Hordeum vulgăre (Barley).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooideae;
Triticeae; Hordeum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zenopus laevis (African clawed frog).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Amphibia, Batrachia, Anura, Mesobatrachia, Pipoidea, Pipidae,
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Viruses, dsDNA viruses, no RNA stage; Baculoviridae;
Nucleopolyhedrovirus.
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                                                                                                                                                                                                                                                                                                                                                             "ENOD40 homologue from barley (Hordeum vulgare).";
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AF542513; AAN28683.1; -.
SEQUENCE 12 AA; 1367 MW; 3EBABFA918EB1733 CRC64;
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01-JUN-2003 (TrEMBLrel. 24, Created)
01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
01-MAR-2003 (TrEMBLrel. 23, Last sequence update) 01-MAR-2003 (TrEMBLrel. 23, Last annotation update) ENOD40-like protein.
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Local Similarity 57.1%;
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Kinase; Transferase.
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Schmidt B., Walter H., Moschik G., Paatz C., Werwein M., Schwingel E.,
Korn K.;
                                                                                                                                                                                                                                                                                                                Gaps
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"Identification of a phylogenetically conserved activin-responsive enhancer in the Zic3 gene."; Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases. EMBL; AF506277; AAP20808.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Singh B., Lin A., Wu Z.-C., Gupta R.S.;
Singh B., Lin A., Wu Z.-C., Gupta R.S.;
"Gene Structure for Adenosine Kinase in Chinese Hamster and Human:
High Frequency Mutants of CHO cells involve deletions of several
introns and exons.";
DNA Cell Biol. 0:0-0(2001).
EMBL, AF318953; AAK07186.1; -.
GO; GO:0004001; F:adenosine kinase activity; IEA.
GO; GO:0016301; F:kinase activity; IEA.
GO; GO:0016740; F:transferase activity; IEA.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
NCBI_TaxID=10090;
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                                                                                                                                                                                                                                             Query Match 2.3%; Score 25; DB 13; Length 13; Best Local Similarity 83.3%; Pred. No. 8e+04; Matches 5; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 25; DB 15; Length 13;
Pred. No. 8e+04;
4; Mismatches 4; Indels
                                                                                                                                                   13 AA; 1569 MW; 819B7A159F531861 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQÜENCE 13 AA; 1390 MW; 805408704623D1AA CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       099PBB;
01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Adenosine kinase (EC 2.7.1.20) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human immunodeficiency virus 1.
Viruses; Retroid viruses; Retroviridae, Lentivirus.
VCBL_TaxID=11676;
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Best Local Similarity 38.5%;
Matches 5; Conservative
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Q99PB
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GenCore version 5.1.6
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OM protein - protein search, using sw model

August 30, 2004, 10:40:09; Search time 35.8378 Seconds (without alignments) 302.211 Million cell updates/sec Run on:

US-09-720-469A-44 1083 1 MKVLIAAALIAGSVFFLLLP......VIIADCGKIEVEKPFAIAKE 208

Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

141681 seqs, 52070155 residues Searched:

585 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 8 Maximum DB seq length: 14

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_42:* Database : Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Description	PS4835 canis famil	P81863 pardachirus	_		_	P28525 hordeum vul	P12509 human immun			_			P27429 squalus aca	P17232 vespa manda			1 kassina		7	P83349 sarcophaga	P56917 rana tempor	9	4	_	5 tabanus	tabanus a	P49325 bacillus th	P83328 oncorhynchu	35 rattus	P01370 tremella me	9prz1 c	P56264 litoria xan	634 8
SUMMARIES	EFTU CANFA	PAP1 PARMA	CA31 LITCI	GER2 HORVU	CA32_LITCI	GER1 HORVU	TAT HV1W2	TAT_HV1Z8	RPCH PANBO	HTF_NAUCI	LPAA PORGI	AL11 CARMA	GONL SQUAC	CRBL_VESMA	CRBL_VESTR	TEMF_RANTE	TKN_KASSE	GONZ_CHICK		FIF1_SARBU	TEMA_RANTE	CRBL_VESOR	FIBB_MANLE	AKH_MELML	AKH_TABAT	HTF TABAT	SLAP_BACTG	RS30 ONCMY	HS9A_RAT	TA13 TREME	BRK_ONCMY	CAER LITXA	TKS1_AEDAE
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TKS2_AEDAE HPB9_RANES	RF1 CONSP UN46 CLOPA LMT2 LOCMI	COXO_RAT	URA7_HUMAN CORZ_PERAM	TKN4_PSEGU ADFB_TENMO	MAST_VESCR
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CA32 LITCI
ID CA32 LITCI
AC P82090;
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P28526;
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Marmoratus).",
Marmoratus).",
J. Biol. Chem. 261:16704-16713(1986).

J. Biol. Chem. 261:16704-16713(1986).

J. Biol. Chem. 261:16704-16713(1986).

J. Biol. Chem. 261:16704-16713(1986).

J. Properties. Forms voltage-dependent, ion-permeable channels in membranes. At high concentration causes cell membrane lysis. Shown to be 5-10 times more toxic, cytolytic and active in membrane pore
                                                                                                                              MEDLINE=87057369; PubMed=3782138;
Lazarovici P., Primor N., Loew L.M.;
"Purification and pore-forming activity of two hydrophobic
polypeptides from the secretion of the Red sea moses sole (Pardachirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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TISSUE=Skin secretion;

MEDLINE=20057701; PubMed=10589099;

Wabnitz P.A., Bowie J.H., Tyler M.J.;

"Caerulein-like peptides from the skin glands of the Australian blue montains tree frog Litoria citropa. Part 1. Sequence determination using electrospray mass spectrometry.";

Rapid Commun. Mass Spectrom. 13:2498-2502(1999).

-!- FUNCTION: Hypotensive neuropeptide (Probable).

-!- SUBCELLULAR LOCATION: Secreted.

-!- TISSUE SPECIFICITY: Skin dorsal glands.

-!- PTM: Isoform 3.1Y4 differs from isoform 3.1 in not being
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-!- SIMILARITY: Belongs to the gastrin/cholecystokinin family.
InterPro; IPR001651; Gastrin.
PROSITE; PS00259; GASTRIN; FALSE_NEG.
Amphibian defense peptide; Hypotensive agent; Amidation; Sulfation;
Pyrrolidone carboxylic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Caerulein 3.1/3.1Y4.
Litoria citropa (Australian blue mountains tree frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Hylidae;
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -!- SUBUNIT: Monomer. In aqueous solution exists as a tetramer.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the pardaxin family.
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Pred. No. 4.6e+03;
1; Mismatches 1; Indels
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10 AA; 1063 MW; D399C36760572DD9 CRC64;
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10DAB7D67861A86B CRC64;
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                                                                                                      TISSUE=Skin secretion;
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NCBI_TaxID=31087;
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(By similarity).
-!- SUBCELLULAR LOCATION: SOLUBLE, MICROSOMAL AND CELL WALL FRACTIONS.
-!- TISSUE SPECIFICITY: Roots and coleoptile. In roots, present in the mature region, but not in the tip. Not detected in leaves.
-!- INDUCTION: Increased by salt stress in roots and decreased by salt stress in coleoptile.
                                                                                                                                                                                                                                                                                                                                                                                                   -!- FUNCTION: May play a role in altering the properties of cell walls during germinative growth.
-!- SUBUNIT: Oligomer (believed to be a pentamer but probably hexamer)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=Skin secretion;
MEDLINE=20057701; PubMed=10589099;
Wabnitz P.A., Bowie J.H., Tyler M.J.;
"Caerulein-like peptides from the skin glands of the Australian blue montains tree frog Litoria citropa. Part 1. Sequence determination using electrospray mass spectrometry.";
                                                                                                                   Hordeum vulgare (Barley).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooldeae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia; Batrachia, Anura, Neobatrachia, Hyloidea, Hylidae,
Pelodryadinae, Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; 0
                                                                                                                                                                                                                                                                                     STRAIN=cv. CM 72; TISSUE=Root;
Hurkman W.J., Tao H.P., Tanaka C.K.;
"Germin-like polypeptides increase in barley roots during salt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 30; DB 1; Length 13;
Pred. No. 6.1e+03;
1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Litoria citropa (Australian blue mountains tree frog)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -!- PTM: Glycosylated.
-!- SIMILARITY: Belongs to the germin family.
InterPro; IPR001929; Germin.
PROSITE; PS00725; GERMIN; PARTIAL.
Apoplast; Cell wall; Glycoprotein; Multigene family.
UNSURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13 AA; 1484 MW; 43FB4A1AA3B7B6D7 CRC64;
                                                                  Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
 13 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE, AND MASS SPECTROMETRY.
                                                                                                                                                                                                                                                                                                                                                           stress.";
Plant Physiol, 97:366-374(1991)
                                        (Rel. 24, Created)
(Rel. 24, Last sequ
(Rel. 40, Last anno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.88;
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STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6; Conservative
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                                                                                                            Germin GS2 (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Caerulein 3.2/3.2Y4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (By similarity)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                  Triticeae, Horde
NCBI_TaxID=4513;
                                  01-DEC-1992
01-DEC-1992
16-OCT-2001
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NON TER
SEQUENCE
                      HV1W2
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-!- FUNCTION: May play a role in altering the properties of cell walls during germinative growth.

-!- SUBUNIT: Oligomer (believed to be a pentamer but probably hexamer) (By similarity).

-!- SUBCELLULAR LOCATION: SOLUBLE, MICROSOMAL AND CELL WALL FRACTIONS.

-!- TISSUE SPECIFICITY: Roots and coleoptile. In roots, present in the mature region, but not in the tip. Not detected in leaves.

-!- INDUCTION: Increased by salt stress in roots and decreased by salt stress in coleoptile.
                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                   Hordeum vulgare (Barley).
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, Pooideae,
        -!- FUNCTION: Hypotensive neuropeptide (Probable).
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Skin dorsal glands.
-!- PTM: Isoform 3.2Y4 differs from isoform 3.2 in not being sulfated.
-!- MASS SPECTROMETRY: MW=1423; METHOD=Electrospray.
-!- SIMILARITY: Belongs to the gastrin/cholecystokinin family.
InterPro; IPR001651; Gastrin.
PROSITE; PS00259; GASTRIN; FALSE NEG.
Amphibian defense peptide; Hypotensive agent; Amidation; Sulfation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE.
STRAIN=cv. CM 72; TISSUE=Root;
Hurkman W.J., Tao H.P., Tanaka C.K.;
"Germin-like polypeptides increase in barley roots during salt
                                                                                                                                                                                                       Score 29; DB 1; Length 11;
Pred. No. 6.2e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.7%; Score 29; DB 1; Length 13; 55.6%; Pred. No. 7.4e+03; Live 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 13;
                                                                                                                                              PYRROLIDONE CARBOXYLIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13 AA; 1470 MW; 43FB588AA3B7B6D7 CRC64;
                                                                                                                                                                      AMIDATION.
10DAB8867861A86B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -!- PTM: Glycosylated.
-!- SIMILARITY: Belongs to the germin family.
InterPro; IPR001929; Germin.
PROSITE; PS00725; GERMIN; PARTIAL.
Apoplast; Cell wall; Glycoprotein; Multigene family.
UNSURE 10 10
Commun. Mass Spectrom. 13:2498-2502(1999)
                                                                                                                                                                                                                                                                                                                                                                   01-DEC-1992 (Rel. 24, Created)
01-DEC-1992 (Rel. 24, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Germin GSL (Fragment).
                                                                                                                                                                                                                                                                                                                                             13 AA.
                                                                                                                                                           SULFATION.
                                                                                                                                                                                                                                                                                                                                             PRT;
                                                                                                                                 Pyrrolidone carboxylic acid. MOD_RES 1
                                                                                                                                                                                                        2.7%;
80.0%;
                                                                                                                                                                      11 11
11 AA; 1363 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 55.0°,
                                                                                                                                                                                                                     Local Similarity 80.0 es 4; Conservative
                                                                                                                                                                                                                                                                                                                                              STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          185 PLKDVIIAD 193
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                                                                                                                                                                                                                                                         125 YGPGW 129
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                                                                                                                                                                                                                                                                                 4 YGTGW 8
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SEQUENCE
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                                                                                                                                                                                   SEQUENCE
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                                                                                                                                                         MOD_RES
MOD_RES
                                                                                                                                                                                                                                                                                                                     LT 6
HORVU
 Rapid
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                                                                                                                                                                                                                                                                                                                                                                                                                 or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            trans-activating responsive sequence (TAR) RNA element and activates transcription initiation and/or elongation from the LTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SUBUNIT: Binds cyclin T1 (By similarity).
SUBCELLULAR LOCATION: Nuclear; nucleolar.
MISCELLANEOUS: ISOLATES WMJ1, WMJ2, AND WMJ3 WERE OBTAINED FROM BLOOD SAMPLES SEQUENTIALLY TAKEN FROM A TWO-YEAR OLD HAITIAN WHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
MEDLINE=86235450; PubMed=3012778;
Hahn B.H., Shaw G.M., Taylor M.E., Redfield R.R., Markham P.D.,
Salahuddin S.Z., Wong-Staal F., Gallo R.C., Parks E.S., Parks W.P.;
"Genetic variation in HTLV-III/LAV over time in patients with AIDS o
at risk for AIDS.";
Science 232:1548-1553(1986).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FUNCTION: Transcriptional regulator that acts by binding to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yourno J., Josephs S.F., Reitz M.S. Jr., Zagury D., Wong-Staal F., Gallo R.C.;
Gallo R.C.;
"Nucleotide sequence analysis of the env gene of a new Zairian isolate of HIV-1.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Transcription regulation; Activator; RNA-binding; Nuclear protein;
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01-OCT-1989 (Rel. 12, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
TAT protein (Transactivating regulatory protein) (Fragment).
                                                                                                                                        TAT protein (Transactivating regulatory protein) (Fragment).
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11681;
                                                                                                                                                                                      Human immunodeficiency virus type 1 (WMJ2 isolate) (HIV-1). Viruses, Retroid viruses, Retroviridae, Lentivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Indels
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14 AA; 1467 MW; 37CC737BFEF67AA8 CRC64;
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75.0%; Pred. No. 8.1e+03;
tive 0; Mismatches 2;
                                                   (Rel. 12, Created)
(Rel. 12, Last sequence update)
(Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14 AA.
  14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WAS PERINATALLY INFECTED BY HER MOTHER.
PRT;
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MEDLINE=88281278; PubMed=3395517;
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HIV; M12507; TAT$WMJ2.
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les 6; Conservative
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STANDARD;
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                                                                                                                                                                                                                                                NCBI_TaxID=11705;
                                                       01-OCT-1989
01-OCT-1989
                                                                                                             28-FEB-2003
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P12511;
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                                                                                                                                                                                                           P10939;
                                                                                                            RESULT 10
HTF NAUCI
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Matches
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                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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01-FEB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Red pigment concentrating hormone (RPCH).
Pandalus borealis (Northern red shrimp).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Bucarida; Decapoda; Pleocyemata; Caridea; Pandaloidea;
AIDS Res. Hum. Retroviruses 4:165-173(1988).
-!- FUNCTION: Transcriptional regulator that acts by binding to the trans-activating responsive sequence (TAR) RNA element and activates transcription initiation and/or elongation from the LTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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-!- FUNCTION: This hormone adapts the animal to light backgrounds by stimulating concentration of the pigment of its red body-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV; J03653; TAT$JY1.
Transcription regulation; Activator; RNA-binding; Nuclear protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=75054965; PubMed=4433569;
Fernlund P.;
"Structure of the red-pigment-concentrating hormone of the shrimp,
                                                                                                                                   promoter.
-!- SUBUNIT: Binds cyclin T1 (By similarity).
-!- SUBCELLULAR LOCATION: Nuclear; nucleolar.
-!- MISCELLANEOUS: THE Z-84 ISOLATE WAS TAKEN FROM A 54 YEAR-OLD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ..
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InterPro; IPR002047; AKH.

PROSITE; PS00256; AKH; 1.

Pigment; Hormone; Amidation; Pyrrolidone carboxylic acid.

MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.

MOD_RES 8 AA; 948 MW; 86786775B9C44736 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.6%; Score 28; DB 1; Length 8; 50.0%; Pred. No. 1.4e+05; live 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chromatophores.
-- SUBCELLULAR LOCATION: Secreted.
!- SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NON_TER 1 1 SEQUENCE 14 AA; 1453 MW; 37CC737BF82D7AA8 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; J03653; AAA44685.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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Best Local Similarity
Matches 3; Conserv
                                                                                                                                                                                                                                                                     ZAIREAN MALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=6703;
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P08939;
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RPCH PANBO
ID RPCH PANBO
DT 01-NOV.
DT 01-NOV.
DT 01-CCE.
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OC EURALIA.
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CC -!- SUI
CC -!- SU
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124 HYGPGW 129

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RA SEQUENCE.

RX MEDLINE=91179584; PubMed=2080017;
RA Veenstra J.A., Camps F.;
RA Veenstra J.A., Camps F.;
RI "Structure of the hypertrehalosemic neuropeptide of the German cockroach, Blattella germanica.";
RI "Structure of the hypertrehalosemic neuropeptides that cockroach, Blattella germanica.";
RL Neuropeptides 15:107-109(1990).
CC -!- FUNCTION: Hypertrehalosemic factors are neuropeptides that cC major carbohydrate in the hemolymph of insects).
CC -!- SUBCELLULAR LOCATION: Secreted. The hemolymph of insects).
CC -!- SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
DR PIR; A26381; A26381.
DR PIR; A60421; A60421.
DR PIR; S08999; S08997.
DR PIR; S08999; S08997.
DR PROSITE; PS00256; AKH; 1.
KW Neuropeptide; Amidation; Pyrrolidone carboxylic acid.
FT MOD_RES 1 1 1 PYRROLIDONE CARBOXY...IC ACID.
FT MOD_RES 1 0 0 AMIDATION.
FT MOD_RES 1 0 AMIDATION.
FORTH MOD_RES 1 0 AMIDATION.
FORT
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MEDLINE=90253659; PubMed=2340112;

Gaede G., Rinehart K.L. Jr.;

"Primary structures of hypertrehalosaemic neuropeptides isolated from

the corpora cardiaca of the cockroaches Leucophaea maderae,

Gromphadorhina portentosa, Blattella germanica and Blatta orientalis

and of the stick insect Extatosoma tiaratum assigned by tandem fast

atom bombardment mass spectrometry.";

Biol. Chem. Hoppe-Seyler 371:345-354(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                         01-JUL-1989 (Rel. 11, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Hypertrehalosaemic hormone (HTH) (Hypertrehalosaemic neuropeptide).
Nauphoeta cinerea (Cinereous cockroach) (Gray cockroach),
Leucophaea maderae (Madeira cockroach),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SPECIES=N.cinerea; TISSUB=Corpora cardiaca;
MEDLINE=87100208; PubMed=3801028;
Gaede G., Rinehart K.L. Jr.;
"Amino acid sequence of a hypertrehalosaemic neuropeptide from the
corpus cardiacum of the cockroach, Nauphoeta cinerea.";
Biochem. Biophys. Res. Commun. 141:774-781(1986).
                                                                                                                                                                                                                                                                                                                                                                                                 Blattella germanica (German cockroach), and Gromphadorina portentosa (Madagascan hissing cockroach).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Orthopteroidea; Dictyoptera; Blattaria; Blaberoidea; Blaberidae; Nauphoeta.
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Pred. No. 6.9e+03;
2; Mismatches 1; Indels
10 AA
PRT;
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STANDARD;
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Best Local Similarity
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-!- FUNCTION: Mast cell degranulating peptide. Induces the chemotaxis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Neoptera; Endopterygota; Hymenoptera; Apocrita; Aculeata; Vespoidea;
Vespidae; Vespinae; Vespa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Squalea; Squalidae; Squalidae; Squalus.
NCBI_TaxID=7797;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Machorina C.S., Feter R.E., Rivier J.E., Sherwood N.M.;

Lovejoy D.A., Fischer W.E., Rivier J.E., Sherwood N.M.;

Nahorniak C.S., Peter R.E., Rivier J.E., Sherwood N.M.;

"Distinct sequence of gonadotropin-releasing hormone (GnRH) in dogfish brain provides insight into GnRH evolution.";

Proc. Natl. Acad. Sci. U.S.A. 89:6373-6377(1992).

-!- FUNCTION: Stimulates the secretion of gonadotropins.

C.!- SUBCELLULAR LOCATION: Secreted.

C.!- SIMILARITY: Belongs to the GnRH family.

R PIR; A46030; A46030.

InterPro; IPR002012; GnRH.

R PROSITE; PS00473; GnRH; 1.

RPOSITE; PS00473; GnRH; 1.

ROD RES 10 10 AMIDATION.

G SEQUENCE 10 AA; 1204 MW; 284B32337871F5A3 CRC64;
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(In) Munekata E. (eds.);
Peptide chemistry 1983, pp.185-190, Protein Research Foundation,
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                                                                                               01-AUG-1992 (Rel. 23, Created)
01-AUG-1992 (Rel. 23, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Gonadoliberin (Gonadotropin-releasing hormone) (GnRH) (LH-RH)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 27; DB 1; Length 10;
Pred. No. 8.4e+03;
3; Mismatches 2; Indels
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13 AA; 1384 MW; 2650402B9DF92338 CRC64;
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01-AUG-1990 (Rel. 15, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Vespid chemotactic peptide M (VESCP-M).
Vespa mandarinia (Hornet).
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                                              10 AA.
                                              PRT;
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                                                                                                                                                                                                                                                   Squalus acanthias (Spiny dogfish)
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Local Similarity 37.5%;
Les 3; Conservative
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                                              STANDARD;
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Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Brain;
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SEQUENCE
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-: FUNCTION: IS ASSOCIATED WITH LIPID A, A PHOSPHORYLATED GLYCOLIPID THAT ANCHORS THE LIPOPOLYSACCHARIDE TO THE OUTER MEMBRANE OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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Eur. J. Biochem. 250:727-734(1997).
-!- FUNCTION: May act as a neurotransmitter or neuromodulator.
-!- SIMILARITY: Belongs to the allatostatin family.
Neuropeptide; Amidation; Multigene family.
MOD RES
9 AMIDATION.
SEQÜENCE 9 AA; 927 MW; 832D79CDCB46DR61 CPC64.
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TISSUE=Cerebral ganglion, and Thoracic ganglion;
MEDLINE=98121193; PubMed=9461295;
Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
Thorpe A.;
                                              15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Lipid-A-associated protein (Fragment).
Porphyromonas gingivalis (Bacteroides gingivalis).
Bacteria; Bacteroidetes; Bacteroides (class); Bacteroidales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Carcinustatin 11.

Carcinus maenas (Common shore crab) (Green crab).

Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;

Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyura;

Eubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 28; DB 1; Length 13;
Pred. No. 9.1e+03;
2; Mismatches 2; Indels
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13 13
13 AA; 1346 MW; 38EA796EAFA63AB7 CRC64;
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30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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13 AA.
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PRT;
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STANDARD;
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LPAA PORGI
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SEQUENCE
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7

Mismatches

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Conservative
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                                                                                                                                                                                                                                                                                                     Kassinin.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Osaka (1983).
-!- FUNCTION: Mast cell degranulating peptide. Induces the chemotaxis
                                                                                                                                                                                                    01-AUG-1990 (Rel. 15, Created)
01-AUG-1990 (Rel. 15, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Vespid chemotactic peptide T (VESCP-T).
Vespa tropica (Hornet).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Hymenoptera; Apocrita; Aculeata; Vespoidea;
Vespidae; Vespinae; Vespa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             temporaria.";
Eur. J. Biochem. 242:788-792(1996).
-!- FUNCTION: Has antibacterial activity against Gram-negative and Gram-positive bacteria.
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Skin.
-!- SIMILARITY: Belongs to the brevinin family.
Amphibian defense peptide; Antibiotic; Amidation.
MOD RES 13 13 AMIDATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rana temporaria (European common frog).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Ranoidea, Ranidae, Rana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Barra D.; "Temporins, antimicrobial peptides from the European red frog Rana temporaria.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yasuhara T., Nakajima T., Erspaer V.;
(In) Sakakibara S. (eds.);
Peptide chemistry 1982, pp.213-218, Protein Research Foundation,
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MEDLINE-97175050; PubMed-9022710;
Simmaco M., Mignogna G., Canofeni S., Miele R., Mangoni M.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 27; DB 1; Length 13;
Pred. No. 1.1e+04;
3; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mast cell degranulation; Chemotaxis; Amidation.
MOD RES 13 13 AMIDATION.
SEOUENCE 13 AA; 1354 MW; 220140365DFE5338 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (Rel. 39, Created)
(Rel. 40, Last sequence update)
(Rel. 42, Last annotation update)
                                                                                                                                                                 13 AA
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Best Local Similarity 44.4
Local 4; Conservative
                                                                                                                                                                 STANDARD;
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             160 VFGKVLEGM 168
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4 IIGKLLSGL 12
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4 ILGKILGGL 12
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TEMF_RANTE
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Best Local Similarity

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H
Gaps
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Squalus acanthias (Spiny dogfish),
Hydrolagus colliei (Spotted ratfish) (Pacific ratfish), and
Clupea pallasii (Pacific herring).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Experientia 33:857-858(1977).
-!- FUNCTION: Tachykinins are active peptides which excite neurons, evoke behavioral responses, are potent vasodilators and secretagogues, and contract (directly or indirectly) many smooth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kassina senegalensis (Senegal running frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Hyperoliidae;
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-!- TISSUE SPECIFICITY: Skin.
-!- SIMILARITY: Belongs to the tachykinin family.
PIR; S07206; S07206.
PDB; 1MYU; 16-OCT-02.
InterPro; IPR002040; Tachy_Neurokinin.
Pfam; PF02202; Tachykinin; 1.
PROSITE; PS00267; TACHYKININ; 1.
Amphibian defense peptide; Tachykinin; Neuropeptide; Amidation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE=Skin secretion;
MEDLINE=77246385; PubMed=891753;
Anastasi A., Montecucchi P.C., Erspamer V., Visser J.;
"Amino acid composition and sequence of kassinin, a tachykinin dodecapeptide from the skin of the African frog Kassina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1;
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01-FEB-1991 (Rel. 17, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Gonadoliberin II (Gonadotropin-releasing hormone II) (GnRH-II)
(IH-RH II) (Luliberin II).
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Pred. No. 1.1e+04;
1; Mismatches 2; Indels
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01-AUG-1988 (Rel. 08, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                         12 AA
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12 AA; 1336 MW;
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4 LIGKVLSGI 12
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Query Match
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P81751;
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MOD_RES
SEQUENCE
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MEDLINE=20114351; PubMed=10650929;
Carolsfeld J., Powell J.F.F., Park M., Fischer W.H., Craig A.G.,
Chang J.P., Rivier J.E., Sherwood N.M.;
"Primary structure and function of three gonadotropin-releasing
hormones, including a novel form, from an ancient teleost, herring.";
Endocrinology 141:505-512(2000).
-!- FUNCTION: Stimulates the secretion of gonadotropins.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the GRRH family.
PIR; B46030; B46030.
PIR; B60066; RHAQ2.
                                                                                                                                                                                            Lovejoy D.A., Fischer W.H., Parker D.B., McRory J.E., Park M., Lance V., Swanson P., Rivier J.E., Sherwood N.M.; "Primary structure of two forms of gonadotropin-releasing hormone from brains of the American alligator (Alligator mississippiensis)."; Regul. Pept. 33:105-116(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                         Lovejoy D.A., Sherwood N.M., Fischer W.H., Jackson B.C., Rivier J.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Primary structure of gonadotropin-releasing hormone from the brain of a holocephalan (ratfish: Hydrolagus colliei)."; Gomp. Endocrinol. 82:152-161(1991).
                                                                                         chicken hypothalamus: evidence that gonadotropin secretion is probably controlled by two distinct gonadotropin-releasing hormones in avian species.";
                                                                  Matsuo H.; "Identification of the second gonadotropin-releasing hormone in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ·.
                                                                                                                                                                                                                                                                                        SPECIES=S.acanthias; TISSUE=Brain; MEDLINE=92335300; PubMed=1631133; Lovejoy D.A., Fischer W.H., Ngamvongchon S., Craig A.G., Nahorniak C.S., Peter R.E., Rivier J.E., Sherwood N.M.; "Distinct sequence of gonadotropin-releasing hormone (GnRH) in dogfish brain provides insight into GnRH evolution."; Proc. Natl. Acad. Sci. U.S.A. 89:6373-6377(1992).
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MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
                       SPECIES=Chicken; TISSUE=Hypothalamus;
MEDLINE=84222059; PubMed=6427779;
Miyamoto K., Hasegawa Y., Nomura M., Igarashi M., Kangawa K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 26; DB 1; Length 10; Pred. No. 1e+04;
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284B2E437871F5A3 CRC64;
                                                                                                                                Proc. Natl. Acad. Sci. U.S.A. 81:3874-3878(1984)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2; Mismatches
                                                                                                                                                                      SPECIES=A.mississippiensis; TISSUE=Brain; MEDLINE=91352338; PubMed=1882082;
                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=91340067; PubMed=1678723;
                                                                                                                                                                                                                                                                                                                                                                                                                 SPECIES=H.colliei; TISSUE=Brain;
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PROSITE; PS00473; GNRH; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KHYGPGW 129
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  [1]
SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE
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                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                             SEQUENCE
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10 AA.

PRT;

STANDARD;

RESULT 19 GON3_ONCKE ID_GON3_ONCKE

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SEQUENCE, AND FUNCTION.
SPECIES=C.pallasi; TISSUE=Brain, and Pituitary;
MEDLINE=20114351; PubMed=10650929;
Carolsfeld J., Powell J.F.F., Park M., Fischer W.H., Craig A.G.,
Chang J.P., Rivier J.E., Sherwood N.M.;
"Primary structure and function of three gonadotropin-releasing
hormones, including a novel form, from an ancient teleost, herring.";
Endocrinology 141:505-512(2000).
-: FUNCTION: Stimulates the secretion of gonadotropins; it stimulates
the secretion of both luteinizing and follicle-stimulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sarcophaga bullata (Grey flesh fly) (Neobellieria bullata).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Oestroidea;
Sarcophagidae; Sarcophaga.
01-FEB-1991 (Rel. 17, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Gonadoliberin III (Gonadotropin-releasing hormone III) (GnRH-III) (LH-RH III) (Luliberin III).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                          Sherwood N., Eiden L., Brownstein M., Spiess J., Rivier J., Vale W.; "Characterization of a teleost gonadotropin-releasing hormone."; Proc. Natl. Acad. Sci. U.S.A. 80:2794-2798(1983).
                                                                                                                                                                                Clupea pallasii (Pacific herring).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Actinopterygii; Neopterygii; Teleostei; Buteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
NCBI_TaxID=8018, 30724;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ġ
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Meeusen T., Mertens I., Clynen E., Baggerman G., Nichols R.,
Nachman R.J., Huybrechts R., De Loof A., Schoofs L.;
"Identification in Drosophila melanogaster of the invertebrate
protein-coupled FMRFamide receptor.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hormone; Amidation; Hypothalamus; Pyrrolidone carboxylic acid.
MOD_RES 1 1 PYRROLIDONE CARBOXXLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 26; DB 1; Length 10;
Pred. No. 1e+04;
3; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      284B3233786B45A3 CRC64;
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FIF1 SARBU

ID FIF1 SARBU

AC P83349;

DT 28-FEB-2003 (Rel. 41, Created)

DT 28-FEB-2003 (Rel. 41, Last sequence update)

DT 28-FEB-2003 (Rel. 41, Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the GnRH family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AMIDATION.
                                                                                                                                                                 (Chum salmon), and
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                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=83195140; PubMed=6341999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10 AA; 1230 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PIR; A21114; A21114.
InterPro; IPR002012; GnRH.
Pfam; PF00446; GnRH; 1.
PROSITE; PS00473; GNRH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 QHWSYGWL 8
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us-09-720-469a-44.aug30.rsp

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Best Local Similarity
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MOD_RES
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FIBE MANLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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01-AUG-1990 (Rel. 15, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Histamine releasing peptide II (HR-II).
Vespa orientalis (Oriental hornet).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Hymenoptera; Apocrita; Aculeata; Vespoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Temporins, antimicrobial peptides from the European red frog Rana temporaria,";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Ranoidea, Ranidae, Rana.
                                                                                                                                                                                                                                                      .;
                                                          -!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the FARP (FMRFamide related peptide)
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Proc. Natl. Acad. Sci. U.S.A. 99:15363-15368(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eur. J. Biochem. 242:788-792(1996),
-!- FUNCTION: Has antibacterial activity against Gram-positive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE-Skin secretion;
MEDLINE-97175050; PubMed-9022710;
Simmaco M., Mignogna G., Canofeni S., Miele R., Mangoni M.L.,
                                                                                                                                                                                                            Length 12;
                                                                                                                                                                                                       Score 26; DB 1; Length 12;
Pred. No. 1.2e+04;
2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.4%; Score 26; DB 1; Length 13; 50.0%; Pred. No. 1.4e+04; ive 3; Mismatches 1; Indels
                                                                                                                                              AMIDATION.
2DC45519C14AB5A7 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13 13 AMIDATION.
13 AA; 1398 MW; 2653612B9DECD408 CRC64;
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-!- TISSUE SPECIFICITY: Skin.
-!- SIMILARITY: Belongs to the brevinin family.
Amphibian defense peptide; Antibiotic; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-MAY-2000 (Rel. 39, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                        13 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rana temporaria (European common frog).
                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
                                                                                                                                                                                                       2.4%;
                                                                                                                                                                 12 AA; 1389 MW;
                                                                                                                                                                                                                                                   4; Conservative
                                                                                                                       Neuropeptide; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                   STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AND SYNTHESIS.
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                                                                                                                                                                                                                                                                                        62 PKTVDNFV 69
                                                                                                                                                                                                                                                                                                                                POPSDNFI 10
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IGRVLSGI 12
                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
Matches 4; Conserv
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                                           junctions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bacteria
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Temporin A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4,
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P17236;
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                                                                                                      family
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SEQUENCE
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SEQUENCE.
                                                                                                                                              MOD RES
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ID CRBL VI
AC P17236
DT 01-AUG
DT 01-AUG
DT 28-FEB
DE Histam
OS Vespa
OC Eukary
OC Neopte
OC Vespid
OX NCBL T
RN [1]
RP SEQUEN
                                                                                                                                                                                                                                                                                                                                                                                                               TEMA_RANTE
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 NO COCCE
SO COCE
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Miroshnikov A.I., Snezhkova L.G., Nazimov I.V., Reshetova O.I., Rozynov B.V., Gushchin I.S.; "Structure and properties of histamine releasing peptides from the venom of Vespa orientalis hornet."; Bioorg. Khim. 7:1467-1477(1981).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    monomers that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=69115139; PubMed=4974768;
Doolittle R.F., Glascow C., Mross G.A.;
"Characterization of fibrinopeptides A and B from a drill (Mandrillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          aggregation.
SUBUNIT: HEXAMER CONTAINING 2 SETS OF 3 NONIDENTICAL CHAINS
(ALPHA, BETA AND GAMMA), LINKED TO EACH OTHER BY DISULFIDE BONDS.
PTM: Conversion of fibrinogen to fibrin is triggered by thrombin, which cleaves fibrinopeptides A and B from alpha and beta chains, and thus exposes the N-terminal polymerization sites responsible for the formation of the soft clot.
                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Cercopithecidae,
Cercopithecinae, Mandrillus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             leucophaeus).";
Biochim. Biophys. Acta 175:217-219(1969).
-!- FUNCTION: Fibrinogen has a double function: yielding monome:
polymerize into fibrin and acting as a cofactor in platelet
                                                                                                                                                                                                                                                                                                                                                        .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9
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01-FBB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Fibrinogen beta chain [Contains: Fibrinopeptide B] (Fragment)
                                                                                                                                                                                                                                                                                                  Score 26; DB 1; Length 14;
Pred. No. 1.5e+04;
6; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 26; DB 1; Length 14;
Pred. No. 1.5e+04;
2; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR002181; Fibrinogen C.
PROSITE; PS00514; FIBRIN AG C DOMAIN; PARTIAL.
Blood coagulation; Plasma; Pyrrolidone carboxylic acid.
PEPTIDE 1 14 FIBRINOPEPTIDE B.
                                                                                                                                                                                                                              AMIDATION.
22015B4A6CEDFD38 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6695B0F11EF72E1B CRC64;
                                                                                                                                               of neutrophils.

PIR, JN0390, JN0390.

Mast cell degranulation; Chemotaxis; Amidation.

MOD RES 14 14 14 AMIDATION.

SEQUENCE 14 AA; 1524 MW; 22015B4A6CEDFD38 C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14 AA.
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                                                                                                                                                                                                                                                                                                      2.4%;
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                                                                                                                                                                                                                                                  14 AA; 1524 MW;
                                                                                                                                                                                                                                                                                                                        30.0%;
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                                                                                                                                                                                                                                                                        STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                  159 VVFGKVLEGM 168
                                                                                                                                                                                                                                                                                                                                                                                                                                               4 LILGKLVKGL 13
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P25423;
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Jaffe H., Raina A.K., Riley C.T., Fraser B.A., Nachman R.J.,
Vogel V.W., Zhang Y.-S., Hayes D.K.;

"Primary structure of two neuropeptide hormones with adipokinetic and
hypotrehalosemic activity isolated from the corpora cardiaca of horse
flies (Diptera).";

L. Proc. Natl. Acad. Sci. U.S.A. 86:8161-8164(1989).

C. -: FUNCTION: This hormone, released from cells in the corpora
cardiaca after the beginning of flight, causes release of
diglycerides from the fat body and then stimulates the flight
muscles to use these diglycerides as an energy source.

C. -: SUBCELLULAR LOCATION: Secreted.

C. -: SUBCELLULAR LOCATION: Secreted.

C. -: SUBCELLULAR LOCATION: Sereted.

C. -: SUBCELLULAR LOCATION: Flight, Pyrrolidone carboxylic acid.

ROSITE: PSO0256; AKH: 1.

RODERES

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MODERES

RAMIDATION.

PRES

RAMIDATION.

SEQÜENCE 8 AA; 949 MW; 86786771A9D1A736 CRC64;
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Neoptera; Endopterygota; Diptera; Brachycera; Tabanomorpha; Tabanidae;
                                                                                                                                                                                                                                                                                                                                                                                                 "A unique charged tyrosine-containing member of the adipokinetic hormone/red-pigment-concentrating hormone peptide family isolated and sequenced from two beetle species.";

Biochem. J. 275:671-677(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SPECIES=P.marginata; TISSUE=Corpora cardiaca;
MEDLINE=92265187; PubMed=1586453;
Gaede G., Lopata A., Kellner R., Rinehart K.L. Jr.;
"Primary structures of neuropeptides isolated from the corpora
cardiaca of various cetonid beetle species determined by
pulsed-liquid phase sequencing and tandem fast atom bombardment mass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        spectrometry.";
Biol. Chem. Hoppe-Seyler 373:133-142(1992).
-!- FUNCTION: This hormone, released from cells in the corpora
-!- FUNCTION: This hormone, released from cells in the corpora
cardiaca after the beginning of flight, causes release of
diglycerides from the fat body and then stimulates the flight
muscles to use these diglycerides as an energy source.
                                                                                                                                                                                                                                                                                                                        SPECIES=M.melolontha, and G.stercorosus; TISSUE=Corpora cardiaca;
MEDLINE=91248100; PubMed=2039445;
                                                                                                                                                                              Eukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota, Neoptera, Endopterygota, Coleoptera, Polyphaga, Scarabaeiformia, Scarabaeidae, Melolonthinae, Melolontha.
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01-FEB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Adipokinetic hormone (AKH) (Dipteran corpora cardiaca factor I)
(DCC I).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ·,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Neuropeptide, Amidation, Flight, Pyrrolidone carboxylic acid.
MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.3%; Score 25; DB 1; Length 8; 50.0%; Pred. No. 1.4e+05; ative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
PIR; A58641; A58641.
PIR; S15422; S15422.
PIR; S21663; S21663.
InterPro; IPR002047; AKH.
PROSITE; PS00256; AKH; 1.
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         01-MAY-1992 (Rel. 22, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
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                                                                                  Adipokinetic hormone (AKH).
Melolontha melolontha (Cockchafer),
Geotrupes stercorosus (Dor beetle), and
Pachnoda marginata (Flower beetle).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AMIDATION
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MEDLINE=90046758; PubMed=2813385;
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Best Local Similarity
Matches 3; Conserv
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P14595;
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Jaffe H., Raina A.K., Riley C.T., Fraser B.A., Nachman R.J.,
Jaffe H., Raina A.K., Riley C.T., Fraser B.A., Nachman R.J.,
Vogel V.W., Zhang Y.-S., Hayes D.K.;
"Primary structure of two neuropeptide hormones with adipokinetic and
hypotrehalosemic activity isolated from the corpora cardiaca of horse
flies (Diptera).";
flies (Diptera).";
From The Card. Sci. U.S.A. 86:8161-8164(1989).
-!- FUNCTION: Hypertrehaloseemic factors are neuropeptides that
elevate the level of trehalose in the hemolymph (trehalose is the
major carbohydrate in the hemolymph of insects).
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
                                                             01-JAN-1990 (Rel. 13, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Hypertrehalosaemic factor (HOTH) (Dipteran corpora cardiaca factor II)
                                                                                                                                                                                                                                     Tabanus atratus (Horse fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Tabanomorpha; Tabanidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Neuropeptide, Amidation; Pyrrolidone carboxylic acid.
MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AMIDATION.
916036786771A9D1 CRC64;
10 AA.
PRT;
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10 AA; 1169 MW;
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InterPro, IPR002047; AKH.
PROSITE; PS00256; AKH; 1.
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STANDARD;
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MEDLINE=907811; PubMed=2592346;
Luckevich M.D., Beveridge T.J.;
"Characterization of a dynamic S layer on Bacillus thuringiensis.";
J. Bacteriol. 171:6656-6667(1989).
-!- FUNCTION: The S-layer is a paracrystalline mono-layered assembly of proteins which coat the surface of bacteria.
-!- SUBCELLULAR LOCATION: CELL WALL. THIS BACTERIUM IS COVERED BY A S-LAYER WITH OBLIQUE (P2) SYMMETRY.
PIR; A60476; A60476.
Cell wall; S-layer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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"A novel antimicrobial function for a ribosomal peptide from rainbow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Actinopterygii, Neopterygii, Teleostei, Euteleostei,
Protacanthopterygii, Salmoniformes, Salmonidae, Oncorhynchus.
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-!- SIMILARITY: Belongs to the S30E family of ribosomal proteins.
Robosomal protein; Antibiotic.
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-!- FUNCTION: Has antibacterial activity against Gram-positive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.3%; Score 25; DB 1; Length 10; 66.7%; Pred. No. 1.3e+04; live 1; Mismatches 1; Indels
                                                                                   01-FEB-1996 (Rel. 33, Created)
01-FEB-1996 (Rel. 33, Last sequence update)
01-FEB-1996 (Rel. 33, Last annotation update)
S-layer protein (Surface layer protein) (Fragment).
Bacillus thuringiensis (subsp. galleriae).
Bacteria, Firmicutes; Bacillales; Bacillaceae; Bacillus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.3%; Score 25; DB 1; Length 11; 71.4%; Pred. No. 1.4e+04;
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10 AA; 1080 MW; 57AECACAB769D1A3 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RS30 ONCMY STANDARD; PRT; 11 AA. P833<u>78</u>; 28-FEB-2003 (Rel. 41, Created) 28-FEB-2003 (Rel. 41, Last sequence update) 10-OCT-2003 (Rel. 42, Last annotation update) 40S ribosomal protein S30 (Fragment).
                                                                 10 AA.
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                                                                 PRT;
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MEDLINE=22142142; PubMed=12147245;
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Matches 5; Conserv
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                              SLAP_BACTG
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RESULT 27
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"Peptide sex hormones inducing conjugation tube formation in compatible mating-type cells of Tremella mesenterica.";
Science 212:1525-1527(1981).
-!- FUNCTION: Tremerogen A-13 is produced by the a mating-type cells and induces formation of conjugation tubes in a mating-type cells. PIR; A01641; JTJG3.
Lipoprotein; Prenylation; Pheromone.
LIPID 13 13 8-farnesyl cysteine.
SEQUENCE 13 AA; 1204 MW; 680304A9697BA864 CRC64;
                                                                                                                           Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota, Fungi, Basidiomycota, Hymenomycetes, Heterobasidiomycetes,
Tremellomycetidae, Tremellales, Tremellaceae, Tremella.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                     STRAIN-Sprague-Dawley; TISSUE-Liver;
MEDLINE=21589773; PubMed=11732320;
Langer T., Fasold H.;
"Isolation and quantification of the heat shock protein 90 alpha and
                                                                                                                                                                                                                                                                                                                                                                                                                                   (BY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (BY
                                                                                                                                                                                                                                                                                                                                         -!- SUBUNIT: Homodimer (By similarity).
-!- SUBCELLULAR LOCATION: Cytoplasmic.
-!- SIMILARITY: Belongs to the heat shock protein 90 family.
InterPro; IPR001404; Hsp90.
PROSITE; PS00298; HSP90; PARTIAL.
Chaperone; ATP-binding; Heat shock; Phosphorylation.
MOD_RES 4 4 PHOSPHORYLATION (BY DS-DNA KINASE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                SIMILARITY).
PHOSPHORYLATION (BY DS-DNA KINASE)
SIMILARITY).
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Pred. No. 1.5e+04;
6; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DE47C322CAB6C1B6 CRC64;
                                                         28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Heat shock protein HSP 90-alpha (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
                               12 AA
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                                                                                                                                                                                                                                                                                 beta isoforms from rat liver.";
Protoplasma 218:54-56(2001).
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12 AA; 1432 MW;
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                              STANDARD;
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ETQTQDQPME 12
                                                                                                                                                                                                                                                                                                                                similarity).
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                              HS9A RAT
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ID TA13_TRE.
P01370;
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                                            P82995;
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TA13 TREME
RESULT 29
HS9A RAT
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1; Indels

1; Mismatches

Conservative

131 SMANAGK 137

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TGEKGFG 79
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les 4; Conserv
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Best Local Similarity
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TGDKFYG
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TKS2_AEDAE
ID _TKS2_AEDAE
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Matches
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Conlon J.M., Olson K.R.;
"Purification of a vasoactive peptide related to lysyl-bradykinin from trout plasma.";
FEBS Lett. 334:75-78(1993).
-!- FUNCTION: SMOOTH MUSCLE CONTRACTION. PROBABLY PLAYS A ROLE FOR THIS SYSTEM IN CARDIOVASCULAR REGULATION IN FISH.
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Plasma.
-!- TISSUE SPECIFICITY: Plasma.
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                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE=Skin secretion;
MEDLINE=97374000; PubMed=9230483;
Steinborner S.T., Waugh R.J., Bowie J.H., Wallace J.C., Tyler M.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Litoria xanthomera (Orange-thighed frog).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Hyloidea, Hylidae,
Pelodryadinae, Litoria.
                                                                                                                                                                                             Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
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        Score 24.5; DB 1; Length 13; Pred. No. 1.8e+04; 2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                          2.2%; Score 24; DB 1; Length 10; llarity 100.0%; Pred. No. 1.5e+04; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                 PIR; S39030; S39030.
Bradykinin; Vasodilator.
SEQUENCE 10 AA; 1193 MW; 33C59075A3786777 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
                                                                                                                                                                   16-OCT-2001 (Rel. 40, Last sequence update) 16-OCT-2001 (Rel. 40, Last annotation update) Lysyl-bradykinin-like.
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                                                                                                                                   10 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Australian tree frog Litoria xanthomera.";
J. Pept. Sci. 3:181-185(1997).
-!- FUNCTION: Hypotensive neuropeptide.
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Skin dorsal glands.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE, AND MASS SPECTROMETRY.
                                                                                                                                                          16-OCT-2001 (Rel. 40, Created)
            2.3%;
Query Match
Best Local Similarity 50.v-
                                                       95 QGGDFTRGDGTG 106
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                                                                           1 EGGG-NRGDPSG 11
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Matches 3; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=79697;
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                                                                                                            RESULT 31
BRK_ONCMY
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MEDLINE = 94105119; PubMed = 8278354;
Champagne D.E., Ribeiro J.M.C.;
Champagne D.E., 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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-!- MASS SPECTROMETRY: MW=1354; METHOD=FAB.
-!- SIMILARITY: Belongs to the gastrin/cholecystokinin family.
InterPro; IPR001651; Gastrin.
PROSITE; PS00259; GASTRIN; 1.
Amphibian defense peptide; Hypotensive agent; Amidation; Sulfation; Pyrrolidone carboxylic acid.
MOD_RES 1 1 SULFATION.
MOD_RES 4 4 SULFATION.
MOD_RES 10 10 AMIDATION.
SEQÜENCE 10 AA; 1290 MW; 99DBF3837861BB5A CRC64;
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01-NOV-1995 (Rel. 32, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Sialokinin II.
Aedes aegypti (Yellowfever mosquito).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea; Aedes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sialokinin I.
Aedes aegypti (Yellowfever mosquito).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea; Aedes.
NCBI_TaxID=7159;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 24; DB 1; Length 10;
Pred. No. 1.5e+04;
1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
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Local Similarity
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UN46 CLOPA
ID UN46 CLOPA
AC P81362;
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                                         "Sialokinin I and II: vasodilatory tachykinins from the yellow fever mosquito Aedes aegypti.";
Proc. Natl. Acad. Sci. U.S.A. 91:138-142(1994).
-!- FUNCTION: Vasodilatory peptide. May activate macrophages at the site of feeding.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the tachykinin family.
                                                                                                                                                                                                                        Gaps
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MEDLINE=90198965; PubMed=2317508;
Simmaco M., de Biase D., Severini C., Aita M., Erspamer G.F.,
Barra D., Bossa F.;
"Purification and characterization of bioactive peptides from skin
                                                                                                                                                                                                                                                                                                                                         01-0CT-1993 (Rel. 27, Created)
01-0CT-1993 (Rel. 27, Last sequence update)
10-0CT-2003 (Rel. 42, Last annotation update)
Hemolytic protein B9 (Fragment).
Rana esculenta (Edible frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Rana.
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                                                                                                                                                                           10 AA; 1146 MW; 3DC77C6B59C33AAB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13 AA; 1402 MW; C6B41A765DF9287D CRC64;
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Amphibian defense peptide; Amidation; Hemolysis.
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(Rel. 41, Last sequence update)
(Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                         13 AA.
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         STRAIN=Rockefeller; TISSUE=Salivary gland; MEDLINE=94105119; PubMed=8278354; Champagne D.E., Ribeiro J.M.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         extracts of Rana esculenta.";
Biochim. Biophys. Acta 1033:318-323(1990).
-!- FUNCTION: Shows hemolytic activity.
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Skin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AMIDATION.
                                                                                                                                                                  AMIDATION
                                                                                                                  PIR; B49581; B49581.

InterPro; IPR002040; Tachy Neurokinin.
PROSITE; PS00267; TACHYKININ; 1.

Tachykinin; Neuropeptide; Amidation.
MOD_RES 10 AMIDATION
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                                                                                                                                                                                                                    4; Conservative
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Best Local Similarity
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2 TGDKFYG 8
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RF1 CONSP
ID RF1 CONSP
AC P58805;
DT 28-FEB-2003 ()
DT 28-FEB-2003 ()
DT 28-FEB-2003 ()
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SEQUENCE
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P32416;
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HPB9_RANES
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Olivera B.M., Heimer de la Cotera E.P.;
"Conorfamide, a Conus venom peptide belonging to the RFamide family of
neuropeptides.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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"Two-dimensional gel electrophoresis separation and N-terminal sequence analysis of proteins from Clostridium pasteurianum W5.";
Electrophoresis 19:802-806(1998).
-! MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED 21 OF THIS UNKNOWN PROTEIN IS: 5.4, ITS MW IS: 38.2 kDa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
15-JUL-1998 (Rel. 36, Last annotation update)
Unknown protein CP 46 from 2D-page (Fragment).
Clostridium pasteurianum.
Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae; Clostridium.
                                                                                                                                                                                                                                                                                                                                                                                        Total of 1401-407 (2002).

-!- FUNCTION: Causes hyperactivity in mice.
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Expressed by the venom duct.
-!- MASS SPECTROMETRY: MW=1454.8; METHOD=Electrospray.
-!- SIMILARITY: Belongs to the FARP (FMRFamide related peptide)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ٦;
Conorfamide-Srl.
Conus spurius (Alphabet cone).
Eukaryota; Metazoa; Mollusca; Gastropoda; Orthogastropoda;
Apogastropoda; Caenogastropoda; Sorbeoconcha; Hypsogastropoda;
Neogastropoda; Conoidea; Conidae; Conus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ..
H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 23.5; DB 1; Length 14; Pred. No. 2.4e+04; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.2%; Score 23.5; DB 1; Length 1 83.3%; Pred. No. 2e+04; Ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOD_RES 12 12 AMIDATION.
SEQUENCE 12 AA; 1456 MW; 2510671E49D772D3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE 14 AA; 1550 MW; 198078F4C0367170 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8 AA.
                                                                                                                                                                                                          SEQUENCE, SYNTHESIS, AND MASS SPECTROMETRY
                                                                                                                                                                                                                                                              MEDLINE=21605839; PubMed=11738233;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE.
STRAIN=W5;
MEDLINE=98291870; PubMed=9629918;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 38
LMT2_LOCMI
ID _LMT2_LOCMI STANDARD;
AC P22396;
DT 01-AUG-1991 (Rel. 19, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         family.
Neurotoxin; Toxin; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.2%;
Local Similarity 38.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       38 VYFDLRIGDEDVG 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       126 GP-GWV 130
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124 HY--GPG 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HYAEGPG 8
|| || 2 HYEEGPG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           URA7_HUMAN
ID _URA7_HUMAN
                                                                                                    THUOB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NON TER
SEQUENCE
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8
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-!- FUNCTION: This protein is one of the nuclear-coded polypeptide chains of cytochrome c oxidase, the terminal oxidase in mitochondrial electron transport.

-!- CATALYTIC ACTIVITY: 4 ferrocytochrome c + O(2) = 4 ferricytochrome
                                                                                                                                                                                                                                                                        -!- FUNCTION: Mediates visceral muscle contractile activity (myotropic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=Wistar; TISSUE=Heart, and Liver;
MEDLINE=95324529; PubMed=7601105;
Schaegger H., Noack H., Halangk W., Brandt U., von Jagow G.;
"Cytochrome-c oxidase in developing rat heart. Enzymic properties and amino-terminal sequences suggest identity of the fetal heart and the adult liver isoform.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rattus norvegicus (Rat).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Cytochrome c oxidase polypeptide VIIc, mitochondrial (EC 1.9.3.1)
(VIIIA) (Fragment).
                                                                                                                                                                                     Schoofs L., Holman G.M., Hayes T.K., Nachman R.J., de Loof A.; "Isolation, identification and synthesis of locustamyotropin II, additional neuropeptide of Locusta migratoria. Member of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              c + 2 H(2)0.
-!- SIMILARITY: Belongs to the cytochrome c oxidase VIIC family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5;
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0
               01-AUG-1991 (Rel. 19, Last annotation update)
Locustamyotropin 2 (LOM-MT-2).
Locusta migratoria (Migratory locust).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridomorpha;
Acridoidea; Acrididae; Oedipodinae; Locusta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 2.1%; Score 23; DB 1; Length 10; Best Local Similarity 71.4%; Pred. No. 1.9e+04; Matches 5; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                           2.1%; Score 23; DB 1; Length 8; 100.0%; Pred. No. 1.4e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10 AA; 1117 MW; 126DE767687B1B0B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                          8 AA; 934 MW; 26341771A9CAA87B CRC64;
                                                                                                                                                                                                                                                                                        activity).
-!- SIMILARITY: Belongs to the pyrokinin family.
InterPro; IPR001484; Pyrokinin.
PROSITE; PS00539; PYROKININ; 1.
Neuropeptide; Amidation; Pyrokinin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10 AA.
    01-AUG-1991 (Rel. 19, Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                           AMIDATION
                                                                                                                                                                                                                                       cephalomyotropic peptide family.";
Insect Biochem. 20:479-484(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oxidoreductase; Mitochondrion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                4; Conservative
                                                                                                                                                                         rissue=Corpora cardiaca;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PIR; S65388; S65388.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97 GDFT 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GDFT 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COX7C OR COX7C1
                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COXO_RAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Loc
Matches
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01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-2001 (Rel. 40, Last annotation update)
Cytochrome c oxidase polypeptide VIIc (EC 1.9.3.1) (Fragment).
Thunnus obesus (Bigeye tuna).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Scombroidei;
Scombridae; Thunnus.
NCBI TaxID=8241;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mitochondrial electron transport.
-!- CATALYTIC ACTIVITY: 4 ferrocytochrome c + 0(2) = 4 ferricytochrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE-Erythrocyte;
Golaz O., Hughes G.J., Frutiger S., Paquet N., Bairoch A.,
Golaz O., Hughes G.J., Tissot J.-D., Appel R.D., Walzer C.,
Basquali C., Sanchez J.-C., Tissot J.-D., Appel R.D., Walzer C.,
Balant L., Hochstrasser D.F.;
Submitted (FEB-1994) to Swiss-Prot.
-!- MISCELLANEOUS: On the 2D-gel the determined pl of this unknown
protein is: 4.77, its MW is: 26 kDa.
SWISS-2DPAGE; P34990; HUMAN.
NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eur. J. Biochem. 248:99-103(1997).
-!- FUNCTION: This protein is one of the nuclear-coded polypeptide chains of cytochrome c oxidase, the terminal oxidase in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "The subunit structure of cytochrome-c oxidase from tuna heart and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-FEB-1994 (Rel. 28, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation update)
Unknown protein from 2D-page of red blood cells (Spot 2D-006HO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Heart, and Liver;
MEDLINE=97454291; PubMed=9310366;
Arnold S., Lee I., Kim M., Song E., Linder D., Lottspeich F.,
Kadenbach B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 2.1%; Score 23; DB 1; Length 10; Best Local Similarity 71.4%; Pred. No. 1.9e+04; Matches 5; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10 AA; 1059 MW; 126DE767687B1DCB CRC64;
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10 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STANDARD;
STANDARD;
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10 AA; 1161 MW; F72D82E9C44B1871 CRC64;

124 HY--GPG 128

1;

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secretagogues, and contract (directly or indirectly) many smooth
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                                                                                                                                                                                                                                                                 4 PDEFFGL 10
                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=7067;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SYNTHESIS.
TISSUE=Head;
                                                                                                                                                                                                                                                                                                                          TENMO
                                                                                                                                                                                            Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        m
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MEDLINE=90287814; PubMed=2356157;
Simmaco M., Severini C., de Biase D., Barra D., Bossa F.,
Roberts J.D., Melchiorri P., Erspamer V.;
"Six novel tachykinin- and bombesin-related peptides from the skin of the Australian frog Pseudophryne guntheri.";
Peptides 11:299-304(1990).
-!- FUNCTION: Tachykinins are active peptides which excite neurons, evoke behavioral responses, are potent vasodilators and
                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DINOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Substance P-like peptide I (PG-SPI).
Pseudophryne guentheri (Guenther's toadlet).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Myobatrachidae; Pseudophryne.
                                                                                                                                                                                                                                                                                                                              the American cockroach.";
FEBS Lett. 250:231-234(1989).
-!- FUNCTION: Cardioactive peptide. Corazonin is probably involved in the physiological regulation of the heart beat.
-!- SUBCELLULAR LOCATION: Secreted.
PIR; S05002; S05002.
                                                                                                                                                                                                                                                                                                                      "Isolation and structure of corazonin, a cardioactive peptide from
                                ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ·;
                                                                                                                                                                                                 Periplaneta americana (American cockroach).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Orthopteroidea; Dictyoptera; Blattaria; Blattoidea;
      Query Match 2.1%; Score 23; DB 1; Length 10; Best Local Similarity 57.1%; Pred. No. 1.9e+04; Matches 4; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 23; DB 1; Length 11; Pred. No. 2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                  PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                          AMIDATION.
C7CFF32D6415AB46 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                    Neuropeptide; Amidation; Pyrrolidone carboxylic acid
                                                                                                                                                       01-OCT-1989 (Rel. 12, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
                                                                                                                                    11 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
                                                                                                                                    PRT;
                                                                                                                                                                                                                                                                                     TISSUE=Corpora cardiaca;
MEDLINE=89325572; PubMed=2753132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                            11 11
11 AA; 1387 MW;
                                                                                                                                  STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STANDARD;
                                                                                                                                                                                                                                       Blattidae; Periplaneta.
NCBI_TaxID=6978;
                                                    111 YGERFPD 117
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                                                                   3 HGENFXD 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               YSRGWTN 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 3; Conserv
                                                                                                                                                                                                                                                                                                          Veenstra J.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 43
TKN4_PSEGU
ID TKN4_PSEGU
AC P42989;
                                                                                                                                  PERAM
                                                                                                                                                                                          Corazonin.
                                                                                                                                                                                                                                                                                                                                                                                                                           MOD_RES
SEQUENCE
                                                                                                                                                                                                                                                                          SEQUENCE
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                                                                                                                                          P11496;
                                                                                                                       CORZ_PERAM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antidiuretic peptide.";
Peptides 24:27-34(2003).
-!- FUNCTION: Inhibitor of fluid secretion by Malpighian tubules. Uses cGMP as second messenger. May function as an antidiuretic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Brain. Appears to be mainly expressed in two pairs of bilaterally symmetrical cells in the protocerebrum.
-!- MASS SPECTROMETRY: MM=1560.33; METHOD=MALDI.
Neuropeptide; Hormone.
SEQUENCE 13 AA; 1562 MW; 0240A4504B8A632B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Κ.Μ.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
C -!- SUBCELLULAR LOCATION: Secreted.
C -!- TISSUE SPECIFICITY: Skin.
C -!- SIMILARITY: Belongs to the tachykinin family.
R PIR; E60409; E60409.
R InterPro; IPR002040; Tachykinin.
R InterPro; IPR008215; Tachykinin.
R Ffam; PF02202; Tachykinin; 1.
R PROSITE; PS00267; Tachykinin; Neuropeptide; Amidation; W Pyrrolidone carboxylic acid.
MOD_RES 1 1 AMIDATION.
C SEQUENCE 11 AA; 1294 MW; 3A247C2CC9CBIAB7 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=22465067; PubMed=12576082;
Eigenheer R.A., Wiehart U.M., Nicolson S.W., Schoofs L., Schegg
Hull J.J., Schooley D.A.;
"Isolation, identification and localization of a second beetle
                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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Neoptera, Endopterygota, Coleoptera, Polyphaga, Cucujiformia,
Tenebrionidae, Tenebrio.
                                                                                                                                                                                                                                                                                                                                                                                                       Score 23; DB 1; Length 11; Pred. No. 2e+04;
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Pred. No. 2.4e+04;
1; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-OCT-2003 (Rel. 42, Created)
10-OCT-2003 (Rel. 42, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Antidiuretic factor B (ADFb).
Tenebrio molitor (Yellow mealworm).
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ID MAST_VESCR

STANDARD;

AC P01516;

DT 21-JUL-1986 (Rel. 01, Created)
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50.0%;
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TISSUE-Venom;

MEDLINE-8428390; PubMed=6206053;

Argiolas A., Pisano J.J.;

"Isolation and characterization of two new peptides, mastoparan C and crabrolin, from the venom of the European hornet, Vespa crabro.";
J. Biol. Chem. 259:10106-10111(1984).

-!- FUNCTION: Mast cell degranulating peptide. Activates G proteins that couple to phospholipase C.

PIR; A01779; QMVHP2.

Mast cell degranulation; Amidation.

MOD RES

14 AA; 1508 MW; 550C0ECA1D6AB1D7 CRC64;
21-JUL-1986 (Rel. 01, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Mastoparan C.
Vespa crabro (European hornet).
Eukaryota, Metazoa, Arthropoda, Hexapoda; Insecta; Pterygota;
Neoptera, Endopterygota, Hymenoptera, Apocrita; Aculeata; Vespoidea;
Vespidae; Vespinae; Vespa.
                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 2.1%; Score 23; DB 1; Length 14; Best Local Similarity 62.5%; Pred. No. 2.6e+04; Matches 5; Conservative 1; Mismatches 2; Indels
                                                                                                                                                          [1]
SEQUENCE.
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Gaps

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Search completed: August 30, 2004, 10:50:26 Job time : 36.8378 secs

1 MKVLLAAA 8

qq ò

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

Run on:

August 30, 2004, 10:41:54; Search time 62.5405 Seconds (without alignments) 319.918 Million cell updates/sec

US-09-720-469A-44 1083 1 MKVLLAAALIAGSVFFLLLP......VIIADCGKIEVEKPFAIAKE 208 Title: Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

283366 segs, 96191526 residues Searched:

1638 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 8 Maximum DB seq length: 14

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

PIR_78:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Result No.	SCORE 331 331 330 239 239	とは のののレントレイ	Lengt	рополого	D TO21 Q135 Q135 TO27 TO27	Description T-cell receptor be T-cell receptor al hypothetical prote Ig heavy chain CRD T-cell receptor be T-cell receptor be t-complex polypept
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14 115 116 118	28 27.5 7.5 7.5		2 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	000000	PH1458 S78766 B56864 A46030 B60769	receptor nal protei idyl-pepti liberin I chain - Pa
010000000000000000000000000000000000000	2222 6666 7222 7222 7222 7225 7225 7225	प्राप्ताय प्राप्त प्राप्त ए बंब बंब बंब बंब बंब वं	4 4 4 8 8 8 9 9 9	000000000	PH0801 S07206 A23695 S33802 PH0747 A58620 PT0554 PT0231	T-cell receptor al kassinin - Senegal myosin heavy chain chaperone, TCP1-re T-cell receptor be adipokinetic hormo adipokinetic hormo T-cell receptor be Ig heavy chain CDR

Trucise
Tracell receptor alpha chain (F15) - mouse (fragment)
Typell receptor alpha chain (F15) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Space: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0796
R;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L.
J. Exp. Med. 174, 1371-1383, 1991
A;Title: T cell receptor genes in a series of class I major histocompatibility complex-reallelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MJD:92078846; PMID:1836010
A;Residues: 1-13 <CAS>
A;Ccoss-references: EMBL:X60901
A;Experimental source: T lymphocyte
C;Keywords: T-cell receptor

RESULT 2

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Gaps

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Score 31; DB 2; Length 13; Pred. No. 9.6e+03; 1; Mismatches 4; Indels

Query Match 2.9%; Best Local Similarity 54.5%; Matches 6; Conservative

ð Dp RESULT 3

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RESULT 1 PT0218 T-cell receptor beta chain V-J region (c) Species: Mus musculus (house mouse) C; Species: Mus musculus (house mouse) C; Date: 31-Dec-1991 #sequence_revision C; Accession: PT0218 R; Nakano, N.; Kikutani, H.; Nishimoto, J. Exp. Med. 173, 1091-1097, 1991 A; Title: T cell receptor V gene usage C A; Reference number: PT0209; MUID:912176 A; Accession: PT0218 A; Molecule type: mRNA A; Residues: 1-11 < NAK> C; Keywords: T-cell receptor	ceptor b : Mus mu : Mus mu on: Frical N.; Kik ed. 173, r cell r ce numbe on: PT02 e type: s: 1-11	SULT 1 2218 2218 2211 receptor beta chain V-J region Species: Mus musculus (house mouse) Date: 31-Dec-1991 #sequence_revision Accession: PT0218 Accession: PT0218 Exp. Med. 173, 1091-1097, 1991 Fitle: T cell receptor V gene usage Reference number: PT0209; MUID:91217 Accession: PT0218 Aolecule type: mRNA Residues: 1-11 <nak></nak>	In V-v (house Lonce, N 197, V gei V gei V gei	d r reginal reginal de din	egion (7 ouse) vision 3 imoto, F usage of :9121763	(7-10-D.3) - mo 31-Dec-1991 #t H.; Kishimoto, of islet beta o 621; PMID:19025	RESULT 1 PT0218 T-cell receptor beta chain V-J region (7-10-D.3) - mouse (fragment C; Species: Mus musculus (house mouse) C; Species: Mus musculus (house mouse) C; Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change 30. C; Accession: PT0218 R; Makano, N.; Kikutani, H.; Nishimoto, H.; Kishimoto, T. J. Exp. Med. 173, 1091-1097, 1991 A; Title: T cell receptor V gene usage of islet beta cell-reactive 'A; Reference number: PT0209; MUID:91217621; PMID:1902501 A; Reference number: mNAA A; Residues: 1-11 caceptor C; Keywords: T-cell receptor	.) -May-199 T cells	7 is not restrict	ict
Query Match Best Local Matches Qy 102	ਕ ਹ	tch al Similarity 50. 5; Conservative 102 GDGTGGKSIY 111 2 GAGQGGNTLY 11	11 11 11	2.9%; 0.0%; ve.ve. 1	Sc. Pr. 2;	ore 31; DB 2; ed. No. 8e+03; Mismatches	; Length ll;	0; Gaps	. 0	

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R;Rommelaere, H.; Van Troys, M.; Gao, Y.; Melki, R.; Cowan, N.J.; Vandekerckhove, J.; Am Proc. Natl. Acad. Sci. U.S.A. 90, 11975-11979, 1993
A;Title: Eukaryotic cytosolic chaperonin contains t-complex polypeptide 1 and seven rela A;Reference number: A49410; MUID:94089752; PMID:7903455
A;Accession: G49410
A;Status: preliminary
A;Molecule type: protein
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A; Title: Structure of the red-pigment-concentrating hormone of the shrimp, Pandalus bore A; Reference number: S07139; MUID:75054965; PMID:4433569
                                                                                        T-cell receptor beta chain V-D-J region (141-1CB) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0588
R;Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Reference number: PT0509; MUID:91277601; PMID:1711558
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N;Alternate names: blanching hormone
C;Species: Pandalus borealis (northern shrimp)
C;Date: 02-Aug-1994 #sequence_revision 05-Aug-1994 #text_change 07-May-1999
C;Accession: A61348; S07139
R;Fernlund, P; J0sefsson, L.
Science 177, 173-175, 1972
A;Title: Crustacean color-change hormone: amino acid sequence and chemical synthesis.
A;Reference number: A61348; MUID:72228738; PMID:5041363
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C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change
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                                                                                                                                                                                                                                                                                                                                                                     A;Molecule type: mRNA
A;Residues: 1-8 <FEE>
A;Experimental source: day 19 fetal thymus, strain BALB/c
C;Keywords: T-cell receptor
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Pred. No. 1.3e+04;
3; Mismatches 3;
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Pred. No. 2.8e+05;
0; Mismatches 1;
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Similarity 45.5%;
5; Conservative 3
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Best Local Similarity 85..
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A;Residues: 1-8 <FER1>
R;Fernlund, P.
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Matches 5; Conserv
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hypothetical protein, 1.3K (rps16 5' region) - Emericella nidulans
C; Species: Emericella nidulans, Aspergillus nidulans
C; Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 09-Sep-1997
C; Accession: JQ1350
R; Bradshaw, R.E.; Pillar, T.M.
Gene 108, 157-162, 1991
A; Title: Isolation and nucleotide sequence of the ribosomal protein S16-encoding gene fr
A; Reference number: JQ1349; MUID:92104500; PMID:1761226
A; Accession: JQ1350
A; Molecule type: DNA
A; Residues: 1-13 <BRA>
A; Residues: 1-13 <BRA>
A; Experimental source: strain R153
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0279
R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0279
A;Molecule type: DNA
A;Residues: 1-8 < YAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
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T-0611 receptor beta chain V-D-J region (140-21) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0725
R;Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Accession: PT0725
A;Status: translation not shown
A;Recession: PT0725
A;Status: translation not shown
A;Residues: 1-8 <FEE>
A;Residues: 1-8 <FEE>
A;Experimental source: newborn thymus, strain BALB/C
C;Keywords: T-cell receptor
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Pred. No. 2.8e+05;
0; Mismatches 1; Indels
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83.3%; Pred. No. 2.8e+05;
ive 0; Mismatches 1; Indels
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Best Local Similarity 83.3
Matches 5, Conservative
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Matches 6; Conserv
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Matches 5; Conserv
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Cispecies: Nauphoeta cinerea (gray cockroach)
Cispecies: Nauphoeta cinerea (gray cockroach)
Cispate: 31-Mar-1988 #sequence_revision 24-Oct-1997 #text_change 31-Oct-1997
Cistate: 31-Mar-1988 #sequence_revision 24-Oct-1997 #text_change 31-Oct-1997
Cistate: 31-Mar-1988 #sequence_revision 24-Oct-1997 #text_change 31-Oct-1997
Cistate: Accession: A26381
Biochem. Biophys. Res. Commun. 141, 774-781, 1986
A;Title: Amino acid sequence of a hypertrehalosaemic neuropeptide from the corpus cardiac A;Reference number: A26381; MUID:87100208; PMID:3801028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Accession: S08998
R;Gaede, G.; Rinehart, K.L.
Biol. Chem. Hoppe-Seyler 371, 345-354, 1990
A;Title: Primary structures of hypertrehalosaemic neuropeptides isolated from the corpore entalis and of the stick insect Extatosoma tiaratum assigned by tandem fast atom bombardn A;Reference number: S08995; MUID:90253659; PMID:2340112
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A;Residues: 1-10 <GAD>
A;Note: the amino-terminal residue forms pyrrolidone carboxylic acid; therefore,
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;10/Modified site: amidated carboxyl end (Thr) #status experimental
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C;Species: Homo sapiens (man)
C;Date: 19-Oct-1995 #sequence_revision 17-Nov-1995 #text_change 05-Nov-1999
C;Accession: S57575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Superfamily: adipokinetic hormone C;Superfamily: adipokinetic hormone, neuropeptide; pyroglutamic acid C;Keywords: amidated carboxyl end; hormone, neuropeptide; pyroglutamic acid F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental F;10/Modified site: amidated carboxyl end (Thr) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypertrehalosemic neuropeptide Bld-HrTH - Madeira cockroach
C;Species: Leucophaea maderae (Madeira cockroach)
C;Date: 30-Jun-1992 #sequence_revision 14-Sep-1994 #text_change 24-Oct-1997
                                                                                                                           F;10/Modified site: amidated carboxyl end (Thr) #status experimental
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50.0%; Pred. No. 1.3e+04;
:ive 2; Mismatches 1; Indels
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A; Molecule type: protein
A; Molecule type: protein
A; Molecule type: protein
A; Molecule type: protein
A; Note: the amino-terminal residue forms pyrrolidone carboxylic acid; therefore, we have
C; Comment: This neuropeptide, isolated from the eyestalks of the shrimp and active in pi
zed pigment-containing cells.
C; Superfamily: adipokinetic hormone
C; Keywords: amidated carboxyl end; blocked amino
F; 1/Modified site: pyrrolidone
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N;Alternate names: Bld-HrTH
C;Species: Blattella germanica (German cockroach)
C;Species: Blattella germanica (German cockroach)
C;Date: 03-Feb-1993 #sequence_revision 03-Feb-1993 #text_change 31-Oct-1997
C;Accession: A60421; S09137
R;Veenstra, J.A.; Camps, F.
Neuropeptides 15, 107-109, 1990
A;Title: Structure of the hypertrehalosemic neuropeptide of the German cockroach, Blatte
A;Reference number: A60421; MUID:91179584; PMID:2080017
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R;Gaede, G.; Rinehart, K.L.
Biol. Chem. Hoppe-Seyler 371, 345-354, 1990
A;Title: Primary structures of hypertrehalosaemic neuropeptides isolated from the corpor entalis and of the stick insect Extatosoma tiaratum assigned by tandem fast atom bombard A;Reference number: S08995; MUID:90253659; PMID:2340112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: protein
A;Residues: 1-10 <VEE>
R;Gaede, G.; Rinehart, K.L.
Biol. Chem. Hoppe-Seyler 371, 345-354, 1990
A;Title: Primary structures of hypertrehalosaemic neuropeptides isolated from the corpor
entalis and of the stick insect Extatosoma tiaratum assigned by tandem fast atom bombard
A;Reference number: S08995; MUID:90253659; PMID:2340112
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A;Residues: 1-10 <GAE>
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
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C.Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;10/Modified site: amidated carboxyl end (Thr) #status experimental
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C;Date: 30-Jun-1992 #sequence_revision 14-Sep-1994 #text_change 24-Oct-1997
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Pred. No. 1.3e+04;
2; Mismatches 1;
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            4; Conservative
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A;Molecule type: protein
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C;Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 17-Mar-1999
C;Accession: PH1458; 226543
R;Casanova, J.L; Martinon, F.; Gournier, H.; Barra, C.; Pamnetier, C.; Regnault, A.; KC
J. Exp. Med. 177; 811-820, 1993
A;Title: T cell receptor selection by and recognition of two class I major histocompatib
A;Reference number: PH1430; MUID:93171821; PMID:8436911
A;Reference number: PH1430; MUID:93171821; PMID:8436911
A;Residues: 1-12 <CAS>
A;Residues: 1-2 <CAS>
A;Residues: 1-12 <AS>
A;Residues: D:: Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Barra, C.; Wid
J. Exp. Med. 176, 439-447, 1992
A;Reference number: S26512; MUID:92364546; PMID:1380061
A;Reference number: S26512; MUID:92364546; PMID:1380061
A;Reference number: EMBL:X67993
A;Residues: 1-12 <CAZ>
A;Cross-references: EMBL:X67993
A;Experimental source: Cytolytic T-lymphocyte, clone Cw3/C44
C;Superfamily: immunoglobulin homology
C;Reywords: receptor; T-cell
R;Burrows, S.R.; Silins, S.L.; Moss, D.J.; Khanna, R.; Misko, I.S.; Argaet, V.P. submitted to the EMBL Data Library, June 1995
A;Description: T cell receptor repertoire for a viral epitope in humans is diversified BA;Reference number: S57494
A;Reference number: S57494
A;Recession: S57575
A;Accession: Pryel minary
A;Molecule type: mRNA
A;Residues: 1-11 < BUR>
A;Cross-references: EMBL: Z49953; NID: 9887510; PIDN: CAA90224.1; PID: 9887511
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tibosomal protein MRP-S28, mitochondrial - bovine (fragment)
C;Species: Bos primigenius taurus (cattle)
C;Bate: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 20-Sep-1999
C;Accession: S78766
R;Graack, H.R.
submitted to the Protein Sequence Database, July 1999
A;Reference number: S78760
A;Accession: S78766
A;Molecule type: protein
A;Residues: 1-13 <GRA>
C;Keywords: mitochondrion
F;1-13/Product: ribosomal protein MRP-S28 (fragment) #status experimental <MAT>
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C;Species: Mus musculus (house mouse)
C;Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 17-Mar-1999
                                                                                                                                                                                                                                                                                                                                       Gaps
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Pred. No. 1.6e+04;
1; Mismatches 2; Indels
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QGSDYTFG 12
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5 GGKLIFG 11
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Best Local Similarity
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Best Local Similarity
Matches 5; Conserv
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PH1458
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dipeptidyl-peptidase IV (EC 3.4.14.5) - bovine (fragment)
C;Species: Bos primigenius taurus (cattle)
C;Species: Bos primigenius taurus (cattle)
C;Date: 05-Jan-1996 #sequence_revision 05-Jan-1996 #text_change 05-Jan-1996
C;Accession: B56864
R;Plakidou-Dymock, S.; McGivan, J.D.
Biochim. Biophys. Acta 1145, 105-112, 1993
Biochim. Biophys. Acta 1145, 105-112, 1993
A;Title: The oligomeric structure of renal aminopeptidase N from bovine brush-border mem A;Reference number: A56864; MUID:93136203; PMID:8093665
A;Accession: B56864
A;Accession: B56864
A;Accession: B56864
A;Molecule type: protein
A;Residues: 1-13 <PLA>
A;Molecule type: protein
A;Residues: 1-13 <PLA>
A;Experimental source: renal brush-border membrane vesicles
C;Keywords: dipeptidylpeptide hydrolase
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NyAlernames: gonadotropin-releasing hormone
C;Species: Squalus acanthias (spiny dogfish)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 16-Dec-1998
C;Accession: A46030
R;Lovejoy, D.A.; Fischer, W.H.; Ngamvongchon, S.; Craig, A.G.; Nahorniak, C.S.; Peter, R
Proc. Natl. Acad. Sci. U.S.A. 89, 6373-6377, 1992
A;Title: Distinct sequence of gonadotropin-releasing hormone (GnRH) in dogfish brain pro
A;Reference number: A46030; MUID:92335300; PMID:1631133
A;Accession: A46030
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-10 <LOV>
C;Keywords: hormone; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
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C;Species: Eptatretus stouti (Pacific hagfish)
C;Date: 03-Jun-1993 #sequence_revision 03-Jun-1993 #text_change 03-Jun-1993
C;Accession: B60769
R;Hanley, P.J.; Seppelt, I.M.; Gooley, A.A.; Hook, J.W.; Raison, R.L.
J. Immunol. 145, 3823-3828, 1990
A;Title: Distinct Ig H chains in a primitive vertebrace, Eptatretus stouti.
A;Reference number: A60769; MUID:91060965; PMID:2123225
  Gaps
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     Indels
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Pred. No. 1.9e+04;
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1;

Gaps

1;

Indels Length

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Accession heavy chain, smooth muscle - chicken (fragment)
C;Species: Gallus gallus (chicken)
C;Species: Gallus gallus (chicken)
C;Species: Gallus gallus (chicken)
C;Date: 04-Oct-1991 #sequence_revision 04-Oct-1991 #text_change 02-Feb-2001
C;Accession: A23695
R;Garabedian, T.E.; Yount, R.G.
J. Biol. Chem. 265, 22547-22553, 1990
A;Title: Direct photoaffinity labeling of gizzard myosin with [3H]uridine diphosphate planate number: A23695; MUID:91093106; PMID:1979981
A;Reference number: A23695
A;Accession: A23695
A;Accession: A23695
A;Accession: A23695
C;Superfamily: myosin heavy chain; myosin motor domain homology
C;Keywords: nucleotide binding; P-loop; smooth muscle
F;2-9/Region: nucleotide-binding motif A (P-loop)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cispecies: Avena sativa (oat)
Cispecies: Oz-nec-1993 #sequence_revision 27-Feb-1997 #text_change 17-Mar-1999
Ciscession: S33802
R;Mummert, E.; Grimm, R.; Speth, V.; Eckerskorn, C.; Schiltz, E.; Gatenby, A.A.; Schaefe:
Nature 363, 644-648, 1993
A;Title: A TCP1-related molecular chaperone from plants refolds phytochrome to its photomaly A;Reference number: S33802
A;Reference number: S33802
A;Accession: S33802
A;Accession: S33802
A;Residues: protein
A;Residues: 1-14 < MUM>
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T-cell receptor beta chain (M1) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0747
R;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L.
                    A;Residues: 1-12 <ANA>
C;Superfamily: unassigned animal peptides
C;Keywords: amidated carboxyl end
F;12/Modified site: amidated carboxyl end (Met) #status experimental
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16.2%; Pred. No. 2.5e+04;
tve 2; Mismatches 2;
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63.6%; Pred. No. 2.1e+04;
Live 1; Mismatches 2;
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Pred. No. 2.3e+04;
2; Mismatches 3;
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Best Local Similarity 53.8%;
Matches 7; Conservative
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Matches 7; Conservative
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Best Local Similarity
Matches 6; Conserva
  A; Molecule type: protein
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Experientia 33, 857-858, 1977
A;Title: Amino acid composition and sequence of kassinin, a tachykinin dodecapeptide fro
A;Reference number: S07206; MUID:77246385; PMID:891753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Accession: PH0801
R;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L.
J. Exp. Med. 174, 1371-1383, 1991
A;Title: T cell receptor genes in a series of class I major histocompatibility complex-rallelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
                                                                                                                                                                                                                                                                                                                                            pyruvate decarboxylase (EC 4.1.1.1) 1 - fungus (Fusarium sporotrichioides) (fragment)

C;Species: Fusarium sporotrichioides
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 23-Mar-2001
C;Accession: PA0096
R;Chow, L.P.; Fukaya, N.; Sugiura, Y.; Ueno, Y.; Tabuchi, K.; Tsugita, A.
submitted to JIPID, October 1994
A;Description: Two dimensional polyacrylamide gel electropheresis of Fusarium sporotrich
A;Reference number: PA0096
A;Molecule type: protein
A;Residues: 1-14 <CHO>
C;Keywords: carbon-carbon lyase; carboxy-lyase
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C;Species: Kassina senegalensis (Senegal running frog)
C;Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 02-Sep-2000
C;Accession: S07206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
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                                                 Score 27; DB 2; Length 11; Pred. No. 1.7e+04; 1; Mismatches 0; Indels
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Pred. No. 2.2e+04;
1; Mismatches 5; Indels
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ilarity 55.6%; Pred. No. 2.2e+04;
Conservative 1; Mismatches 3;
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                                                     2.5%;
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Best Local Similarity 50.0%;
Matches 6; Conservative
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A;Residues: 1-14 <CAS>
A;Cross-references: EMBL:X60909
                          36 VKVYFDLRIGDE 47
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Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                 90 KDFMIQ 95
                                                                                                                                                                                                                       EDFMIQ 9
A;Residues: 1-11 <HAN>
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3; Indels

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Query Match
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Wighteric hormone - damselfly (Pseudagrion inconspicuum)

Wighternete names: Psi-AKH

C;Species: Pseudagrion inconspicuum

C;Date: 19-Mar-1997 #sequence_revision 31-Oct-1997 #text_change 07-May-1999

C;Accession: S55310

R;Janssens, M.P.E.; Kellner, R.; Gaede, G.

R;Janssens, M.P.E.; Kellner, R.; Magner, R.; Gaede, G.

R;Janssens, M.; Gaede, G.

R;Janssens, 
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C;Species: Ischnura senegalensis
C;Date: 28-Oct-1997 #sequence_revision 31-Oct-1997 #text_change 07-May-1999
C;Accession: A58620
R;Janssens, M.P.E.; Kellner, R.; Gaede, G.
Biochem. J. 302, 539-543, 1994
A;Title: A novel adipokinetic octapeptide found in the damselflies Pseudagrion inconspic A;Reference number: S55310; MUID:94379987; PMID:8093008
A;Reference number: S55310; MUID:94379987; PMID:8093008
A;Reference number: S55310; MUID:94379987; PMID:8093008
C;Rossion: A58620
A;Residues: 1-8 cJAN>
C;Superfamily: adipokinetic hormone
C;Superfamily: adipokinetic hormone
C;Superfamily: adipokinetic acid (Gln) #status experimental
F;1/Modified site: amidated carboxyl end (Trp) #status experimental
                                             series of class I major histocompatibility complex-
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50.0%; Pred. No. 2.8e+05;
live 2; Mismatches 1; Indels
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                   A;Title: T cell receptor genes in a series of class I malalle exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
A;Accession: PH0747
A;Molecule type: mRNA
A;Residues: 1-14 <CAS>
A;Cross-references: EMBL:X60838
A;Experimental source: T lymphocyte
C;Keywords: T-cell receptor
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Best Local Similarity 50.0%; Pred. No. 2.5e+04;
Matches 6; Conservative 2; Mismatches 1;
Med. 174, 1371-1383, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                104 GTGGKSIYGERF 115
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Best Local Similarity 50.0
Matches 3; Conservative
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Best Local Similarity 50,0
Matches 3; Conservative
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3 NFTPGW 8
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3 NFTPGW 8
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Quadoliberin II - American alligator
NyAlternate names: gonadotropin-releasing hormone II
C;Species: Alligator mississippiensis (American alligator)
C;Date: 31-Mar-1993 #sequence_revision 31-Mar-1993 #text_change 18-Mar-1997
C;Accession: B60066
R;Lovejoy, D.A.; Fischer, W.H.; Parker, D.B.; McRory, J.E.; Park, M.; Lance, V.; Swanson Regul. Pept. 33, 105-116, 1991
A;Title: Primary structure of two forms of gonadotropin-releasing hormone from brains of A;Reference number: A60066; MUID:91352338; PMID:1882082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Species Homo sapiens (man) (line) (
T-cell receptor beta chain V-D-J region (126-1G) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0554
R;Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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A; Molecule type: protein
A; Residues: 1-10 <LOV>
C; Superfamily: gonadoliberin
C; Keywords: amidated carboxyl end; hormone; hypothalamus; pyroglutamic acid
F; 1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F; 10/Modified site: amidated carboxyl end (Gly) #status experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Accession: PT0554
A; Status: translation not shown
A; Molecule type: mRNA
A; Residues: 1-8 <FEE>
A; Experimental source: day 18 fetal thymus, strain BALB/cC; Keywords: T-cell receptor
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50.0%; Pred. No. 2.8e+05;
iive 3; Mismatches 1;
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A;Residues: 1-9 <YAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
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| ASGDEGLG
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DB 1; Length 10;

Score 26;

2.4%;

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Treell receptor beta chain (clone Cw3/HLA1C8) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 13-Jan-1995 #sequence_revision 17-Apr-1998 #text_change 17-Mar-1999
C;Accession: S26544
R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Barra, C.; Widn J. Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T cell receptor: A;Reference number: S26512; MUID: 92364546; PMID: 1380061
A;Reference number: Cabs>
A;Accession: S26544
A;Residues: 1-12 <CAs>
A;Accession: Cabs>
A;Cross-references: EMBL:X67994
A;Experimental source: cytolytic T-lymphocyte, clone Cw3/HLA1C8
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T-cell receptor gamma chain (5a.3) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 07-May-1999
C;Accession: C38887
R;Whetsell, M.; Mosley, R.L.; Whetsell, L.; Schaefer, F.V.; Miller, K.S.; Klein, J.R. Mol. Cell. Biol. 11, 5902-5909, 1991
A;Title: Rearrangement and junctional-site sequence analyses of T-cell receptor gamma A;Reference number: A41946; MUID:92049316; PMID:1658619
A;Accession: C38887
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S65730
hemoglobin, extracellular, component - earthworm (Lumbricus terrestris) (fragment)
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C;Date: 06-Dec-1996 #sequence_revision 13-Mar-1997 #text_change 13-Mar-1997
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                                                                                                                                                                                                                                                                        Gaps
                 A;Title: Characterization of a teleost gonadotropin-releasing hormone. A;Reference number: A21114; MUID:83195140; PMID:6341999
A;Accession: A21114
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-10 <SHE>
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                                                                                                                                                                                        2.4%; Score 26; DB 2; Length 10; 37.5%; Pred. No. 1.9e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 26; DB 2; ]
Pred. No. 2.1e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.4%;
Local Similarity 57.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5; Conservative
                                                                                                                                                                                                                     Query Match
Best Local Similarity 37.5
Matches 3; Conservative
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                                                                                                                                                                                                                                                                                                                                123 KHYGPGWV 130
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Matches 5, Conserva
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N;Alternate names: gonadotropin-releasing hormone
C;Species: Squalus acanthias (spiny dogfish)
C;Date: O6-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 20-Jan-2003
C;Date: O6-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 20-Jan-2003
C;Accession: B46030
R;Lovejoy, D.A.; Fischer, W.H.; Ngamvongchon, S.; Craig, A.G.; Nahorniak, C.S.; Peter, R
Proc. Natl. Acad. Sci. U.S.A. 89, 6373-6377, 1992
A;Title: Distinct sequence of gonadotropin-releasing hormone (GnRH) in dogfish brain prod A;Reference number: A46030; MUID:92335300; PMID:1631133
                                                                                                                                                                                                                                                                                                                                                                                                                 C; Accession: A61126
R; Lovejoy, D.A.; Sherwood, N.M.; Fischer, W.H.; Jackson, B.C.; Rivier, J.E.; Lee, T. Gen. Comp. Endocrinol. 82, 152-161, 1991
A; Title: Primary structure of gonadotropin-releasing hormone from the brain of a holocep A; Reference number: A61126; MUID: 91340067; PMID: 1678723
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C;Species: Oncorhynchus keta (chum salmon)
C;Date: 10-Aug-1990 #sequence_revision 10-Aug-1990 #text_change 18-Jun-1993
C;Accession: A21114
R;Sherwood, N.; Eiden, L.; Brownstein, M.; Spiess, J.; Rivier, J.; Vale, W.
Proc. Natl. Acad. Sci. U.S.A. 80, 2794-2798, 1983
                                                                                                                                                                                                                                                                                                                                N,Alternate names: gonadotropin-releasing hormone
C;Species: Hydrolagus colliei (spotted ratfish)
C;Date: 26-May-1994 #sequence_revision 26-May-1994 #text_change 18-Mar-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Molecule type: protein
A;Residues: 1-10 <LOV>
A;Experimental source: brain
C;Superfamily: gonadoliberin
C;Keywords: amidated carboxyl end; brain; hormone; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;10/Modified site: amidated carboxyl end (Gly) #status experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: protein
A;Residues: 1-10 <LOV>
C;Superfamily: gonadoliberin
C;Keywords: hormone; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.4%; Score 26; DB 1; Length 10; 42.9%; Pred. No. 1.9e+04; tive 2; Mismatches 2; Indels
                                                        Indels
Pred. No. 1.9e+04;
                                                   2; Mismatches
                          42.98;
                                                                                                                                                                                                                                                                                                   gonadoliberin - spotted ratfish
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Best Local Similarity 42.9
Matches 3; Conservative
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Best Local Similarity 42.9
Matches 3; Conservative
                                                          Conservative
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                               Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Accession: A61126
                                                        3;
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                                                        Matches
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Accession: S15422 MulD:91248100; PMID:2039445

A; Reference number: S15422; MulD:91248100; PMID:91248100; PMID
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A33995
adipokinetic hormone - black horse fly
C;Species: Tabanus atratus (black horse fly)
C;Species: Tabanus atratus (black horse fly)
C;Date: 23-Mar-1990 #sequence_revision 23-Mar-1990 #text_change 31-Oct-1997
C;Accession: A33995
R;Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Nachman, R.J.; Vogel, V.W.; Zhang, Proc. Natl. Acad. Sci. U.S.A. 86, 8161-8164, 1989
A;Title: Primary structure of two neuropeptide hormones with adipokinetic and hypotrehal A;Reference number: A33995; MUID:90046758; PMID:2813385
A;Accession: A33995
A;Molecule type: protein
                                                                                                                                                                                                                                                                                                                                                                                                           T-cell receptor alpha chain (K1 V-alpha-4.3) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0795
K;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L.
J. Exp. Med. 174, 1371-1383, 1991
A;Title: T cceptor genes in a series of class I major histocompatibility complex-sallelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
A;Accession: PH0795
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                     Indels
                        Ä
                        Mismatches
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                     į 9
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A;Cross-references: EMBL:X60900
                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4; Conservative
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                                                                                                   159 VVFGXVLEGM 168
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4 LILGXLVKGL 13
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Matches 4; Conserval
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Best Local Similarity
Matches 3; Conserv
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S15422
                 Matches
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C; Species: Vespa orientalis (oriental hornet)
C; Cpecies: Vespa orientalis
R; Miroshnikov, A.I.; Snezhkova, L.G.; Nazimov, I.V.; Reshetova, O.I.; Rozinov, B.V.; Gus
Bioorie : Jula 
C;Accession: S65730
R;Fushitani, K.; Higashiyama, K.; Asao, M.; Hosokawa, K.
Biochim. Biophys. Acta 1292, 273-280, 1996
A;Title: Characterization of the constituent polypeptides of the extracellular hemoglobi
A;Reference number: S65721; MUID:96176855; PMID:8597573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Spate: 02-May-1994 #sequence_revision 18-Nov-1994 #text_change 16-Aug-1996
C;Accession: C53275
R;Ayadi, H.; Marche, P.N.; Cazenave, P.A.
Immunogenetics 34, 201-207, 1991
A;Title: Evolution of the rabbit immunoglobulin kappa chain genes.
A;Reference number: A53275; MUID:91372868; PMID:1909995
A;Accession: C53275
A;Accession: C53275
A;Accession: C53275
A;Actus: preliminary
A;Molecule type: DNA
A;Residues: 1-13 < AYA>
A;Note: sequence extracted from NCBI backbone (NCBIN:56069, NCBIP:56164)
C;Comment: This J3 segment may not be functional because of substitutions in the 7 mer C;Keywords: heterotetramer; immunoglobulin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   histamine-releasing peptide II - oriental hornet
                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 83.5
-has 5; Conservative
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Best Local Similarity 83.3
Matches 5; Conservative
                                                                                                                                                                                                                         A;Accession: S65730
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-12 <FUS>
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Best Local Similarity
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C; Accession: B33995
R; Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Nachman, R.J.; Vogel, V.W.; Zhang, J
Proc. Natl. Acad. Sci. U.S.A. 86, 8161-8164, 1989
A; Title: Primary structure of two neuropeptide hormones with adipokinetic and hypotrehalc A; Reference number: A33995; MUID: 90046758; PMID: 2813385
A; Accession: B33995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Residues: 1-10 <LUC>
C;Comment: The S-layer, or surface array, is the outermost component of several archaebad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Molecule type: protein
A;Residues: 1-10 <JAF>
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamic
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status predicted
F;10/Modified site: amidated carboxyl end (Tyr) #status predicted
                 J. Exp. Med. 174, 115-124, 1991
A; Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A; Reference number: PT0509; MUID:91277601; PMID:1711558
A; Accession: PT0595
A; Status: translation not shown
A; Molecule type: mRNA
A; Residues: 1-8 <FEE>
A; Essidues: 1-8 <FEE>
A; Experimental source: newborn thymus, strain BALB/c
C; Keywords: T-cell receptor
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C;Species: Bacillus thuringiensis
C;Date: 20-Feb-1993 #sequence_revision 20-Feb-1993 #text_change 12-Mar-1993
C;Accession: A60476
R;Luckevich, M.D.; Beveridge, T.J.
J. Bacteriol. 171, 6656-6667, 1989
A;Title: Characterization of a dynamic S layer on Bacillus thuringiensis.
A;Reference number: A60476; MUID:90078111; PMID:2592346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hypotrehalosemic hormone - black horse fly
C;Species: Tabanus atratus (black horse fly)
C;Date: 23-Mar-1990 #sequence_revision 23-Mar-1990 #text_change 31-Oct-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.3%; Score 25; DB 2; Length 10; 60.0%; Pred. No. 2.3e+04; tive 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                             Length 8;
                                                                                                                                                                                                                                                                                           2.3%; Score 25; DB 2; I
80.0%; Pred. No. 2.8e+05;
tive 1; Mismatches 0;
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Best Local Similarity 80.00,
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4 FTPGW 8
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R; Feeney, A.J.
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                       C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status predicted
F;8/Modified site: amidated carboxyl end (Trp) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C.Species: Gentrupes stercorosus (dor beetle)
C;Species: Geotrupes stercorosus (dor beetle)
C;Date: 28-Oct-1997 #sequence_revision 31-Oct-1997 #text_change 31-Oct-1997
C;Accession: A58641
R;Gaede, G.
Biochem. J. 275, 671-677, 1991
A;Title: A unique charged tyrosine-containing member of the adipokinetic hormone/ red-pi
A;Reference number: S15422; MUID:91248100; PMID:2039445
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R;Gaede, G.; Lopata, A.; Kellner, R.; Rinehart, K.L.
Biol. Chem. Hoppe-Seyler 373, 133-142, 1992
A;Title: Primary structures of neuropeptides isolated from the corpora cardiaca of vario
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;8/Modified site: amidated carboxyl end (Trp) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       neuropeptide - flower beetle (Pachnoda marginata)
C;Species: Pachnoda marginata
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 24-Jul-1997
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Best Local Similarity 60.00
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A,Status: preliminary
A,Molecule type: protein
A,Residues: 1-8 <GAE>
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A;Molecule type: protein
A;Residues: 1-8 <BIO>
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Matches 3; Conserv
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  A;Residues: 1-8 <JAF>
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August 30, 2004, 10:49:45; Search time 254.378 Seconds (without alignments) 257.253 Million cell updates/sec
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| cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*
| cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 14
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Perfect score:
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Description	Sequence 52, Appl	Sequence 347, App	Sequence 85, Appl	Sequence 2, Appli	Sequence 84, Appl	Sequence 1, Appli	Sequence 9, Appli	Sequence 7, Appli	P4	Sequence 242, App	Sequence 168, App	Sequence 118, App	Seguence 142, App	Sequence 66, Appl	Sequence 101, App
SUMMARIES	US-10-394-980-52	US-10-014-340-347	US-10-447-161-85	US-10-788-016-2	US-10-447-161-84	US-10-788-016-1	US-10-788-016-9	US-09-972-656-7	US-09-873-637-9	US-09-876-904A-242	US-10-038-612-168	US-10-264-309-118	US-10-356-257-142	US-10-300-694A-66	US-10-247-839-101
DB	15	12	16	16	16	16	16	10	σ	10	13	16	16	14	12
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% Query Match	7.7	5.0	4.6	4.6	4.5	4.5	4.3	3.3	3,3	3.3	3.3	3.2	3.2	3.2	3.2
Score	83	54	20	50	49	49	47	36	36	36	36	35	35	35	35
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ALIGNMENTS

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NAME/KEY: MISC_FEATURE
OTHER INFORMATION: part of CYPB_HUMAN (peptidyl-prolyl cis-trans isomerase
               Sequence 52, Application US/10394980
; Sequence 52, Application US/10394980
; Publication No. US20040005633A1
; GENERAL INFORMATION:
; APPLICANT: Vandekerckhove, Joel
; APPLICANT: Vandekerckhove, Joel
; TITLE OF INVENTION: METHODS AND APPARATUS FOR GEL-FREE QUALITATIVE AND
; TITLE OF INVENTION: QUANTITATIVE PROTEOME ANALYSIS, AND USES THEREFOR
; FILE REFERENCE: UBV-001
; CURRENT RAPLICATION NUMBER: US/10/394,980
; CURRENT RAPLICATION NUMBER: DCT/EP02/03368
; PRIOR APPLICATION NUMBER: US60/278,171
; PRIOR APPLICATION NUMBER: US60/318,749
; PRIOR PILING DATE: 2001-03-22
; PRIOR APPLICATION NUMBER: US60/318,749
; PRIOR PILING DATE: 2001-09-12
; PRIOR APPLICATION NUMBER: US60/323,999
; NUMBER OF SEQ ID NOS: 473
; SOFTWARE: PatentIn Version 3.1
; SEQ ID NO 52
; LENGTH: 14

TVDE: DRA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
US-10-394-980-52
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Gaps .. 0 7.7%; Score 83; DB 15; Length 14; 100.0%; Pred. No. 0.031; tive 0; Mismatches 0; Indels Conservative Best Local Similarity Matches 14; Conserv

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FEATURE:
CTHER INFORMATION: Peptide consisting of 9 amino acid residues from the 91st residue
CTHER INFORMATION: to the 99th residue of cyclophillin B
US-10-788-016-2
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TITLE OF INVENTION: Mutant Fibronectin and Tumor Metastasis
FILE REFERENCE: HO-P02484US1
CURRENT APPLICATION NUMBER: US/10/447,161
CURRENT FILING DATE: 2003-05-28
PRIOR APPLICATION NUMBER: 60/383,530
PRIOR FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 148
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                     4.6%; Score 50; DB 16; Length 9; 100.0%; Pred. No. 1.2e+06; vative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 1.2e+06;
Matches 9; Conservative 0; Mismatches 0;
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TITLE OF INVENTION: Desensitizers
FILE REFERENCE: 3190-049
CURRENT APPLICATION NUMBER: US/10/788,016
CURRENT FILING DATE: 2004-02-26
PRIOR APPLICATION NUMBER: PCT/JP02/08641
PRIOR FILING DATE: 2002-08-28
PRIOR FILING DATE: 2001-08-29
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.2
SEQ ID NO 1
PRIOR APPLICATION NUMBER: JP P2001-260046
PRIOR FILING DATE: 2001-08-29
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.2
SEQ ID NO 2
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTHER INFORMATION: Synthetic Peptide US-10-447-161-84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 84, Application US/10447161; Publication No. US20040023314A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-788-016-1; Sequence 1, Application US/10788016; Publication No. US20040141992A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100...
                                                                                                                                                                                                                                                                                                                                                                                91 DFMIQGGDF 99
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                                                                                                                               TYPE: PRT
ORGANISM: Artificial
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US-10-447-161-84
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                                                                                                                                               APPLICANT: Herath, et al TITLE OF INVENTION: Nucleic Acid Molecules, Polypeptides and Uses Therefor, Including TITLE OF INVENTION: Diagnosis and Treatment of Alzheimer's Disease FILE REFERENCE: 9195-078
CURRENT APPLICATION NUMBER: US/10/014,340
CURRENT FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 823
SOFTWARE: Patentin version 3.0
SEQ ID NO 347
LENGTH: 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Wang, Rong-fu
TITLE OF INVENTION: Mutant Fibronectin and Tumor Metastasis
FILE REFERENCE: HO-P02484US1
CURRENT APPLICATION NUMBER: US/10/447,161
CURRENT FILING DATE: 2003-05-28
PRIOR APPLICATION NUMBER: 60/383,530
PRIOR FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 148
SOFTWARE: Patentin version 3.1
SEQ ID NO 85
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.6%; Score 50; DB 16; Length 9; 100.0%; Pred. No. 1.2e+06; tive 0; Mismatches 0; Indels
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TITLE OF INVENTION: Desensitizers
FILE REFERENCE: 3190-049
CURRENT APPLICATION NUMBER: US/10/788,016
CURRENT FILING DATE: 2004-02-26
PRIOR APPLICATION NUMBER: PCT/JP02/08641
PRIOR FILING DATE: 2002-08-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Synthetic Peptide US-10-447-161-85
                                                                                  Sequence 347, Application US/10014340 Publication No. US20030064411A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-447-161-85; Sequence 85, Application US/10447161; Publication No. US20040023314A1; GENERAL INFORMATION:
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US-10-788-016-2
; Sequence 2, Application US/10788016
; Publication No. US20040141992A1
; GENERAL INFORMATION:
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Best Local Similarity 100.
Matches 9; Conservative
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                                                              US-10-014-340-347
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Sequence 242, Application US/09876904A
; Sequence 242, Application US/09876904A
; Publication No. US20030072794A1
; GENERAL INFORMATION:
; APPLICANT: BOULKAS, TENI
; TITLE OF INVENTION: ENCAPSULATION OF PLASMID DNA (LIPOGENES TM) AND THERAPEUTIC
; TITLE OF INVENTION: CONJUGATES INTO TARGETED LIPOSOME COMPLEXES
; TITLE OF INVENTION: CONJUGATES INTO TARGETED LIPOSOME COMPLEXES
; FILE REFERENCE: TB-2002.00
; CURRENT APPLICATION NUMBER: US 60/210,925
; PRIOR APPLICATION NUMBER: US 60/210,925
; PRIOR FILING DATE: 2000-06-09
; NUMBER OF SEQ ID NOS: 629
; SOFTWARE: PATENTIN Ver. 2.1
; SEQ ID NO 242
                                                                                                                                        US-09-873-637-9
; Sequence 9, Application US/09873637
; Patent No. US20020061543A1
; GENERAL INFORMATION:
; APPLICANT: Ross, Jeffrey
; TITLE OF INVENTION: (CRD-BP) AND ITS NUCLEIC ACID SEQUENCE
; TITLE OF INVENTION: (SP) 95131
; CURRENT APPLICATION NUMBER: US/09/873,637
; UNMBER OF SEQ ID NOS: 46
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
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Best Local Similarity 58.8%; Pred. No. 4.2e+03;
Matches 10; Conservative 1; Mismatches 2;
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ORGANISM: Unknown Organism
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102 GDGTGGKSIY 111
                                              2 GDNLGGKSLH 11
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US-09-876-904A-242
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TYPE: PRT
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    ; FEATURE:
; OTHER INFORMATION: Peptide consisting of 9 amino acid residues from the 84th residue
; OTHER INFORMATION: to the 92nd residue of cyclophillin B
US-10-788-016-1
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Publication No. US2003009647A1
GENERAL INFORMATION:
APPLICANT: Tsai, Mei-Mei
TITLE OF INVENTION: Fully Human Antibody Fab Fragments with Human Interferon-Gamma;
FILE REFERENCE: A-799
CURRENT APPLICATION NUMBER: US/09/972,656
CURRENT FILING DATE: 2001-10-05
NUMBER OF SEQ ID NOS: 135
SOFTWARE: Patentin version 3.0
SEQ ID NO 7
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                                                                                                                        Length 9;
                                                                                                                                                                        0; Indels
                                                                                                                      Query Match
4.5%; Score 49; DB 16; I
Best Local Similarity 100.0%; Pred. No. 1.2e+06;
Matches 9; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                       US-10-788-016-9

Sequence 9, Application US/10788016

Publication No. US20040141992A1

GENERAL INFORMATION:

APPLICANT: ITOH, KYOGO

TITLE OF INVENTION: Desensitizers

FILE REFERENCE: 3190-049

CURRENT FILING DATE: 2004-02-26

PRIOR APPLICATION NUMBER: PCT/JP02/08641

PRIOR FILING DATE: 2002-08-28

PRIOR FILING DATE: 2002-08-28

PRIOR FILING DATE: 2001-08-29

NUMBER OF SEQ ID NOS: 9

SOFTWARE: Patentin version 3.2

SEQ ID NO 9
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ORGANISM: Artificial
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TYPE: PRT
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APPLICANT: Ben-Sasson, Shmuel A

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RESULT 15
US-10-247-839-101
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APPLICANT: FAMEKH, RAJESH B.
APPLICANT: PAREKH, RAJESH B.
APPLICANT: PAREKH, RAJESH B.
APPLICANT: POTTER, DAVID M.
APPLICANT: ROHLFF, CHRISTIAN
APPLICANT: SILBER, B. MICHABEL
APPLICANT: SILBER, B. MICHABEL
APPLICANT: SUNDER, PETER J.
APPLICANT: SUNDERLAND, P. TREY
APPLICANT: SUNDERLAND, P. TREY
APPLICANT: WHITE, W. FROST
APPLICANT: WHITE, W. FROST
APPLICANT: WHITE, W. FROST
APPLICANT: WHITE, W. FROST
APPLICANT: WILLIAMS, STEPHEN A.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES, POLYPEPTIDES AND USES THEREFOR,
TITLE OF INVENTION: INCLUDING DIAGNOSIS AND TREATMENT OF ALZHEIMER'S DISEASE
FILE REFERENCE: POA-002.01
CURRENT APPLICATION NUMBER: US/10/264,309
CURRENT APPLICATION NUMBER: C60/326,708
FRIOR APPLICATION NUMBER: 60/326,708
FRIOR APPLICATION NUMBER: 2001-10-03
NUMBER OF SEQ ID NOS: 491
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TITLE OF INVENTION: Short Peptides Which Selectively
TITLE OF INVENTION: Modulate the Activity of Protein Kinases
FILE REFERENCE: 1242.1029-000 (CMCC-679)
CURRENT APPLICATION NUMBER: US/10/038,612
CURRENT FILING DATE: 2002-01-08
PRIOR APPLICATION NUMBER: US 09/161,094
PRIOR FILING DATE: 1998-09-25
NUMBER OF SEQ ID NOS: 172
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 168
LENGTH: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 3.3%; Score 36; DB 13; Length 13; Best Local Similarity 66.7%; Pred. No. 4.2e+03; Matches 6; Conservative 2; Mismatches 1; Indels
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Pred. No. 1.2e+06;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-264-309-118; Application US/10264309; Sequence 118, Application US/10264309; Publication No. US20040022794A1; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                         TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: DURHAM, L. KATHRYN
APPLICANT: FRIEDMAN, DAVID L.
APPLICANT: HERATH, HERATH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 3.28
Best Local Similarity 77.8
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                               ; LOCATION: (0)...(13); CTHER INFORMATION: TrkB US-10-038-612-168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        119 NFKLKHYGP 127
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: AMIDATION LOCATION: (0)
                                                                                                                                                                                                                                                                                                                                          NAME/KEY: MYRISTATE
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Pred. No. 5.4e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Polycarbonate-binding peptide 66 US-10-300-694A-66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 66, Application US/10300694A; Sequence 66, Application VS/10300694A; Publication No. US20030185870A1; GENERAL INFORMATION:
APPLICANT: Duke University
APPLICANT: Kenan, Daniel J.
APPLICANT: Walsh, Elisabeth B.
APPLICANT: Middleton, Crystan
TITLE OF INVENTION: INTERFACIAL BIOMATERIALS
FILE REFERENCE: 180/13/2;
CURRENT FILING DATE: 2003-05-07
PRIOR APPLICATION NUMBER: US 60/331,843
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 117
SOFTWARE: Patentin version 3.2
SEQ ID NO 66
TABLICATION OF THE OF SEQ ID NO 66
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                                                             APPLICANT: LAZARIOS, ROBERT A.
APPLICANT: MAUN, HENRY R.
TITLE OF INVENTION: FVIIa Antagonists
FILE REFERENCE: P1950R1
CURRENT APPLICATION NUMBER: US/10/356,257
CURRENT APPLICATION NUMBER: US 60/355,420
PRIOR APPLICATION NUMBER: US 60/355,420
PRIOR FILING DATE: 2002-02-06
NUMBER OF SEQ ID NOS: 355
SEQ ID NO 142
                                                                                                                                                                                                                                                                                                                                                                                           CTHER INFORMATION: sequence is synthesized US-10-356-257-142
// Sequence 142, Application US/10356257
// Publication No. US20040087767A1
// GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial sequence
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3.2%;
Best Local Similarity 41.7%;
Matches 5; Conservative
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Best Local Similarity 77.0
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Sequence 209, Application US/10014340
; Sequence 209, Application US/10014340
; Publication No. US20030064411A1
; GENERAL INFORMATION:
; APPLICANT: Herath, et al
; TITLE OF INVENTION: Diagnosis and Treatment of Alzheimer's Disease
; TITLE OF INVENTION: Diagnosis and Treatment of Alzheimer's Disease
; FILE REFERENCE: 9195-078
; CURRENT APPLICATION NUMBER: US/10/014,340
; CURRENT FILING DATE: 2001-12-10
; NUMBER OF SEQ ID NOS: 823
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 209
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Publication No. US20020106325A1

GENERAL INFORMATION:

APPLICANT: Carpenter, Jr., Alan P.

TITLE OF INVENTION: RECEPTOR TARGETED IMAGING OF CARDIAC PERFUSION AND A VITRONECTIN FILE REFERENCE: BMS-2201

CURRENT APPLICATION NUMBER: US/09/995,388

CURRENT FILING DATE: 2001-11-27

PRIOR PPLING DATE: 2000-11-27

NUMBER OF SEQ ID NOS: 49

SOFTWARE: Patentin version 3.2
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            APPLICANT: Waterfield, Michael Derek

TITLE OF INVENTION: Proteins, Genes, and Their Use for

TITLE OF INVENTION: Diagnosis and Treatment of Breast Cancer;

FILE REFERENCE: 2543-1-024

CURRENT APPLICATION NUMBER: US/09/988,493

CURRENT FILING DATE: 2002-05-21

PRIOR APPLICATION NUMBER: PCT/GB01/01219

PRIOR FILING DATE: 2000-03-20

PRIOR FILING DATE: 2000-03-20

PRIOR FILING DATE: 2000-03-20

PRIOR FILING DATE: 2000-02-24

NUMBER OF SEQ ID NOS: 308

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 293
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Best Local Similarity 77.8
Matches 7; Conservative
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Best Local Similarity 77.8
Matches 7; Conservative
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US-10-014-340-209
                                                                                                                                                                                                                                                                                                                                                                                                                                          ; ORGANISM: homo sapien
US-09-988-493-293
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US-09-995-388-8
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ADDRESSE: Marshall O'Toole Gerstein Murray and Borun STREET: 6300 Sears Tower 233 South Wacker Drive CITY: Chicago
                                                                                                                                                                                                                      Multivalent and Multispecific Binding Proteins, Their Manufacture and Use
                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 60606-6402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (BPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/247,839
FILING DATE: 20-Sep-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 3.2%; Score 35; DB 12; Length 14; Best Local Similarity 50.0%; Pred. No. 6e+03; Matches 7; Conservative 2; Mismatches 5; Indels
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Publication No. US20030064419A1
GENERAL INFORMATION:
APPLICANT: Herath, Herath Mudiyanselage Athula Chandrasiri
APPLICANT: O'Hare, Michael John
APPLICANT: Page, Martin John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/146,979
FILING DATE: <Unknown>
APPLICATION NUMBER: US 08/448,418
FILING DATE: 14-MAY-1996
APPLICATION NUMBER: PCT/GB93/02492
FILING DATE: 03-DEC-1993
APPLICATION NUMBER: GB 9225453.1
FILING DATE: 04-DEC-1992
APPLICATION NUMBER: GB 9300816.7
FILING DATE: 16-JAN-1993
APPLICATION NUMBER: EP 93303614.7
FILING DATE: 10-MAY-1993
APPLICATION NUMBER: GB 9319969.3
FILING DATE: 22-SEP-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: David W. Clough
REGISTRATION NUMBER: 36,107
REFERENCE/DOCKET NUMBER: 28111/32651
INFORMATION FOR SEQ ID NO: 101:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide linker
SEQUENCE DESCRIPTION: SEQ ID NO: 101:
Holliger, Kaspar-Philipp
Griffiths, Andrew D
Hoogenboom, Hendricus RJM
Malmqvist, Magnus
                                                                                                                                                                      Prospero, Terence D
Winter, Gregory P
TITLE OF INVENTION: Multivalent
                                                                                                  Marks, James D
McGuinness, Brian T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 14 amino acids
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                                                                                                                                                Pope, Anthony R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96 GGDFTRGDGTGGKS 109
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GGGGSGGGGGGGSS 14
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                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                  STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 16
US-09-988-493-293
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Query Match
Best Local Similarity 70.0
Matches 7; Conservative
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Best Local Similarity 70.0
Matches 7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 7; Conserv
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US-10-193-477-146

j Sequence 146, Application US/10193477

j Publication No. US2030195163A1

j GENERAL INFORMATION:

j TILLS OF INVENTION:

TILLE OF INVENTION: DELYCINE RICH REPEATS AND IMMUNOLOGOBULIN FOLDS, BGS2, 3, AND 4,

TILLE OF INVENTION: THEREOF

TILLE OF INVENTION TOWNER: US/10/193,477

CURRENT APPLICATION NUMBER: US 60/304,888

PRIOR APPLICATION NUMBER: US 60/372,147

PRIOR APPLICATION NUMBER: US 60/372,147

PRIOR FILLING DATE: 2002-04-12

NUMBER OF SEQ ID NOS: 229

SOFTWARE: PATENTIN VERSION 3.1

SEQ ID NO 146

LENGTH: 12
                                                                                                                                                                                                                                                                                                                          RESULT 19
US-09-995-388-5
; Sequence 5, Application US/09995388
; Publication Wo. US20020106325A1
; GENERAL INFORMATION:
; APPLICANT: Carpenter, Jr., Alan P.
; TITLE OF INVENTION: RECEPTOR TARGETED IMAGING AGENT
; TITLE OF INVENTION: RECEPTOR TARGETED IMAGING AGENT
; TITLE OF INVENTION: WINDER: US/09/995,388
; CURRENT APPLICATION NUMBER: US 60/253,324
; PRIOR APPLICATION NUMBER: US 60/253,324
; RIADR APPLICATION NUMBER: US 60/253,324
; NUMBER OF SEQ ID NOS: 49
; SOFTWARE: Patentin version 3.2
; LENGTH: 12
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                                                                                                                                                                3.1%; Score 34; DB 12; Length 11; 85.7%; Pred. No. 5.5e+03; tive 0; Mismatches 1; Indels
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                                                                                                   ; OTHER INFORMATION: Synthetic Construct US-09-995-388-8
                                                          ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                              Query Match
Best Local Similarity 85.7
Matches 6; Conservative
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CRGANISM: Mus musculus
US-10-193-477-146
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Best Local Similarity
Matches 6; Conserv
SEQ ID NO 8
LENGTH: 11
TYPE: PRT
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Sequence 169 application US/10193477

Publication No. US20030195163A1

GENERAL INFORMATION:

APPLICANT: Bristol-Myers Squibb Company

TITLE OF INVENTION: LEUCINE RICH REPEATS AND IMMUNOLOGOBULIN FOLDS, BGS2, 3, AND 4,

TITLE OF INVENTION: THEREOF

FILE REFERENCE: D0153 NP

CURRENT APPLICATION NUMBER: US/10/193,477

CURRENT APPLICATION NUMBER: US 60/304,888

PRIOR FILING DATE: 2001-07-11

PRIOR FILING DATE: 2002-04-12

NUMBER OF SEQ ID NOS: 229

SOFTWARE: Patentin version 3.1

SEQUENCE: SEQUENCE: SEQUENCE: D0168
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US-10-017-193-8

; Sequence 8, Application US/10017193

; Publication No. US20030113478A1

; GENERAL INFORMATION:

; APPLICANT: Dang, Mai Huong

; APPLICANT: Chiu, Phillip

; TITLE OF INVENTION: Surface Coating Method and Coated Device

; FILE REFERENCE: 52200-8010

; CURRENT APPLICATION NUMBER: US/10/017,193

; CURRENT FILING DATE: 2001-12-12

; NUMBER OF SEQ ID NOS: 10

; SEQ ID NO 8

; LENGTH: 13
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Score 34; DB 14; Length 12; Pred. No. 6.2e+03; 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 34; DB 14; Length 12;
Pred. No. 6.2e+03;
1; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 32.5; DB 14;
Pred. No. 1e+04;
2; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
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OTHER INFORMATION: Mutant sequence substituted in place of IGE Fc(epsilon)3 fragment US-09-925-179-50
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Sequence 27, Application US/10223172A

Publication No. US2003013844A1

GENERAL INFORMATION:

APPLICANT: Myriad Genetics, Inc.

APPLICANT: Zavitz, Kenton

APPLICANT: Wettstein, Daniel Albert

APPLICANT: Worham, Scott

APPLICANT: Morham, Scott

APPLICANT: Hobden, Adrian

TITLE OF INVENTION: COMPOSITION AND METHOD FOR TREATING HIV INFECTION

FILE REFERENCE: 5003.01

CURRENT APPLICATION NUMBER: US/10/223,172A

CURRENT FILING DATE: 2002-08-19

PRIOR FILING DATE: 2001-08-18

NUMBER OF SEQ ID NOS: 42

SOFTWARE: PatentIn version 3.1
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                                                                                                                                     US-09-925-179-50

i Sequence 50, Application US/09925179

j Publication No. US20030044858A1

i GENERAL INFORMATION:

APPLICANT: Jardieu, Paula M.

APPLICANT: Presta, Leonard G.

TITLE OF INVENTION: Anti-IGE Antibodies (as amended)

FILE REFERENCE: P0718P2CIDICUS

CURRENT APPLICATION NUMBER: US/09/925,179

CURRENT APPLICATION NUMBER: US 08/466,163

PRIOR FILING DATE: 1995-06-06

PRIOR FILING DATE: 1995-03-15

PRIOR FILING DATE: 1995-03-15

PRIOR FILING DATE: 1992-08-14

PRIOR FILING DATE: 1992-08-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.(
Matches 5; Conservative
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4 WLDGK 8
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PRIOR FILING DATE: 1994-01-26

PRIOR FILING DATE: 1994-01-26

PRIOR FILING DATE: 1992-08-14

PRIOR FILING DATE: 1992-08-07

PRIOR FILING DATE: 1992-08-05

PRIOR FILING DATE: 1992-08-14

PRIOR FILING DATE: 1992-08-14

PRIOR FILING DATE: 1992-08-14

PRIOR FILING DATE: 1992-08-14

PRIOR FILING DATE: 1992-08-05

PRIOR FILING DATE: 1992-08-05

PRIOR FILING DATE: 1992-08-05

PRIOR FILING DATE: 1991-08-14

PRIOR FILING DATE: 1991-08-14

PRIOR FILING DATE: 1991-08-14

NUMBER OF SEQ ID NOS: 64

SEQ ID NOS: 64
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US-09-802-096-50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; FEATURE:
; OTHER INFORMATION: Mutant sequence substituted in place of IGE Fc(epsilon)3 fragment
US-09-802-077-50
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                                                                                                              APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Method of Treating Allergic Disorders (as amended)
FILE REFERENCE: P0718P2C2US
CURRENT APPLICATION NUMBER: US/09/802,077
CURRENT FILING DATE: 2001-03-08
PRIOR FILING DATE: 1995-03-15
PRIOR FILING DATE: 1995-03-15
PRIOR FILING DATE: 1994-01-26
PRIOR FILING DATE: 1994-01-26
PRIOR FILING DATE: 1992-08-14
PRIOR FILING DATE: 1992-08-14
PRIOR FILING DATE: 1992-08-14
PRIOR FILING DATE: 1991-08-14
PRIOR FILING DATE: 1991-08-14
NUMBER OF SEQ ID NOS: 64
SEQ ID NO 50
LENGTH: 9
LENGTH: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 3.0%; Score 32; DB 9; Length 9; Best Local Similarity 100.0%; Pred. No. 1.2e+06; Matches 5; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 1.2e+06;
Matches 5; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 50, Application US/09802096 Patent No. US20010038839A1
                                Sequence 50, Application US/09802077 Patent No. US20010033842A1
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                                                                                       GENERAL INFORMATION:
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153 WLDGK 157

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RESULT 29
US-09-802-077-46
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Sequence 17, Application US/09802077;
Patent No. US20010033842A1
GENERAL INFORMATION:
APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Method of Treating Allergic Disorders (as amended);
FILE REFERENCE: P0718P2C2US
CURRENT APPLICATION NUMBER: US/09/802,077
CURRENT APPLICATION NUMBER: US 08/405,617
PRIOR APPLICATION NUMBER: US 08/185,899
PRIOR FILING DATE: 1994-01-26
PRIOR FILING DATE: 1994-01-26
PRIOR FILING DATE: 1992-08-14
PRIOR FILING DATE: 1992-08-14
PRIOR FILING DATE: 1992-08-14
PRIOR FILING DATE: 1992-08-14
PRIOR FILING DATE: 1991-08-14
NUMBER OF SEQ ID NOS: 64
SEQ ID NO 17
LENGTH: 10
                                                                                                                                                                                                              APPLICANT: Morham, Scott
APPLICANT: Morham, Scott
APPLICANT: Zavitz, Kenton
APPLICANT: Zavitz, Kenton
APPLICANT: Adrian
TITLE OF INVENTION: Composition and Method for Treating Viral Infection
FILE REFERENCE: 5004.01
CURRENT APPLICATION NUMBER: US/10/224,999A
CURRENT FILING DATE: 2003-03
PRIOR FILING DATE: 2001-08-20
NUMBER OF SEQ ID NOS: 3484
SOFTWARE: Patentin version 3.1
SEQ ID NO 27
LENGTH: 9
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8e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Modified HIV Peptide US-10-224-999A-27
                                                                                                                                           ; Sequence 27, Application US/10224999A; Publication No. US20030171318A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
20 PGPSAADEK 28
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                                    1 PGPSAPPEE 9
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Best Local Similarity
Matches 5; Conserva
                                                                                                   RESULT 27
US-10-224-999A-27
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US-09-802-077-17
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153 WLDGK 157

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; FEATURE:
; OTHER INFORMATION: Mutant sequence substituted in place of IgE Fc(epsilon)3 fragment
US-09-802-077-46
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Patent No. US20010038839A1

GENERAL INFORMATION:

APPLICANT: Jardieu, Paula M.

APPLICANT: Dardieu, Paula M.

APPLICANT: Presta, Leonard G.

TITLE OF INVENTION: Method of Preventing the Onset of Allergic Disorders (as amended FILE REFERENCE: P0718P2C3US

CURRENT APPLICATION NUMBER: US/09/802,096

CURRENT FILING DATE: 1995-03-15

PRIOR PILING DATE: 1995-03-15

PRIOR PLING DATE: 1995-08-14

PRIOR APPLICATION NUMBER: US 08/405,617

PRIOR APPLICATION NUMBER: US 07/879,495

PRIOR APPLICATION NUMBER: US 07/879,495

PRIOR APPLICATION NUMBER: US 07/879,495

PRIOR PILING DATE: 1992-08-14

PRIOR PILING DATE: 1991-08-14

PRIOR PILING DATE: 1991-08-14

PRIOR PILING DATE: 1991-08-14

PRIOR PILING DATE: 1991-08-14

NUMBER OF SEQ ID NOS: 64

SEQ DD NO 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                         GENERAL INFORMATION:

APPLICANT: Jardieu, Paula M.
APPLICANT: Jardieu, Paula M.
APPLICANT: Jardieu, Leonard G.
TITLE OF INVENTION: Method of Treating Allergic Disorders (as amended)
FILE REFERENCE: P0718P2C2US
CURRENT APPLICATION NUMBER: US/09/802,077
CURRENT FILING DATE: 2001-03-08
PRIOR APPLICATION NUMBER: US 08/185,899
PRIOR FILING DATE: 1994-01-26
PRIOR FILING DATE: 1994-01-26
PRIOR FILING DATE: 1992-08-14
PRIOR FILING DATE: 1992-05-07
PRIOR FILING DATE: 1992-05-07
PRIOR FILING DATE: 1991-08-14
PRIOR FILING DATE: 1991-08-14
PRIOR FILING DATE: 1991-08-14
PRIOR SPELICATION NUMBER: US 07/744,768
PRIOR FILING DATE: 1991-08-14
NUMBER OF SEQ ID NOS: 64
SEQ ID NO 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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; Sequence 46, Application US/09802077; Patent No. US20010033842A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.
Matches 5; Conservative
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CRGANISM: Homo sapiens
US-09-802-096-17
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Best Local Similarity
Matches 5; Conserv
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; OTHER INFORMATION: Mutant sequence substituted in place of IgE Fc(epsilon)3 fragment US-09-925-179-46
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US-09-755-630A-33

; Sequence 33, Application US/09755630A
; Publication No. US20030194399A1
; Publication No. US20030194399A1
; GENERAL INFORMATION:
; APPLICANT: ALIBHAI, MURTAZA F.
; APPLICANT: SAMPSON, HUGH A.
; APPLICANT: ASTWOOD, JAMES D.
; APPLICANT: MCWHERTER, CHARLES A.
; TITLE OF INVENTION: PREPARATION OF DEALLERGENIZED PROTEINS AND PERMUTEINS
; FILE REFERENCE: 11899.0217.NPUS00 (MOBT217)
; CURRENT APPLICATION NUMBER: US/09/755,630A
; CURRENT APPLICATION NUMBER: US 60/174,669
; PRIOR APPLICATION NUMBER: US 60/174,669
; NUMBER OF SEQ ID NOS: 293
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 33
; LENGTH: 10

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                                                                     3.0%; Score 32; DB 10; Length 10; 100.0%; Pred. No. 8e+03; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 46, Application US/09925179;
Sequence 46, Application US/09925179;
Publication No. US20030044858A1
GENERAL INFORMATION:
APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Anti-1gE Antibodies (as amended);
FILE REFERENCE: P0718P2C1D1C1US
CURRENT APPLICATION NUMBER: US/09/925,179
CURRENT APPLICATION NUMBER: US 08/466,163
PRIOR FILING DATE: 1995-06-06
PRIOR PELING DATE: 1995-06-06
PRIOR FILING DATE: 1995-08-15
PRIOR PELING DATE: 1995-08-14
PRIOR PELING DATE: 1992-08-14
PRIOR FILING DATE: 1991-08-14
PRIOR FILING DATE: 1991-08-14
NUMBER OF SEQ ID NOS: 68
SEQ ID NO 46
LENGTH: 10
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ORGANISM: Artificial sequence
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 5; Conservative
                                                                                                                                                       5; Conservative
                                                                        Query Match
Best Local Similarity
Matches 5; Conserv
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US-09-925-179-46
US-09-925-179-17
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APPLICANT: Jardieu M.
APPLICANT: Dardieu Paula M.
TITLE OF INVENTION: Method of Preventing the Onset of Allergic Disorders (as amended FILE OF INVENTION: Method of Preventing the Onset of Allergic Disorders (as amended FILE REFERENCE: P0718P2C3US
CURRENT APPLICATION NUMBER: US/09/802,036
CURRENT APPLICATION NUMBER: US 08/405,617
PRIOR PILING DATE: 1995-03-15
PRIOR PILING DATE: 1994-01-26
PRIOR FILING DATE: 1994-01-26
PRIOR PILING DATE: 1992-08-14
PRIOR FILING DATE: 1992-08-14
PRIOR FILING DATE: 1992-08-07
PRIOR FILING DATE: 1991-08-14
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US-09-802-096-46
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Publication No. US20030044858A1
GENERAL INFORMATION:
APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Anti-IGE Antibodies (as amended)
FILE REFERENCE: P0718P2C1D1C1US
CURRENT APPLICATION NUMBER: US/09/925,179
CURRENT FILING DATE: 2001-08-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT FILING DATE: 2001-08-08

PRIOR APPLICATION NUMBER: US 08/466,163

PRIOR FILING DATE: 1995-06-06

PRIOR FILING DATE: 1995-03-15

PRIOR FILING DATE: 1995-03-15

PRIOR PPLICATION NUMBER: US 08/185,899

PRIOR PPLICATION NUMBER: PCT/US92/06860

PRIOR PPLICATION NUMBER: PCT/US92/06860

PRIOR FILING DATE: 1992-08-14

PRIOR FILING DATE: 1992-06-14

PRIOR FILING DATE: 1992-05-07

PRIOR PILING DATE: 1992-05-07

PRIOR FILING DATE: 1991-08-14

NUMBER OF SEQ ID NOS: 68

LENGTH: 10

LENGTH: 10
                                                                                                                                                                                                                                                                  US-09-802-096-46; Sequence 46, Application US/09802096; Patent No. US20010038839A1; GENERAL INFORMATION:
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ORGANISM: Homo sapiens
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                                            WLDGK 157
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US-10-224-999A-33
; Sequence 33, Application US/10224999A
; Publication No. US2003017131BA1
; GENERAL INFORMATION:
; APPLICANT: Myriad Genetics, Inc.
; APPLICANT: Morham, Scott
; APPLICANT: Application and Method for Treating Viral Infection
; TITLE OF INVENTION: Composition and Method for Treating Viral Infection
; TITLE OF INVENTION: Composition and Method for Treating Viral Infection
; FILE REFERENCE: 5004.01
; CURRENT APPLICATION NUMBER: US/10/224,999A
; CURRENT FILING DATE: 2001-08-20
; PRIOR FILING DATE: 2001-08-20
; NUMBER OF SEQ ID NOS: 3484
; SOFTWARE: PatentIn Version 3.1
; SEQ ID NO 33
; LENGTH: 10
                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Zavitz Carton
APPLICANT: Wettstein, Daniel Albert
APPLICANT: Wettstein, Daniel Albert
APPLICANT: Morham, Scott
APPLICANT: Hobden, Adrian
TITLE OF INVENTION: COMPOSITION AND METHOD FOR TREATING HIV INFECTION
FILE REFERENCE: 5003.01
CURRENT APPLICATION NUMBER: US/10/223,172A
CURRENT FILING DATE: 2002-08-19
PRIOR FILING DATE: 2001-08-18
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin version 3.1
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                                                                                                   Query Match 3.0%; Score 32; DB 10; Length 10; Best Local Similarity 75.0%; Pred. No. 8e+03; Matches 6; Conservative 1; Mismatches 1; Indels
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                      ; OTHER INFORMATION: Synthetic polypeptide US-09-755-630A-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Modified HIV Peptide US-10-224-999A-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                           Sequence 33, Application US/10223172A Publication No. US20030138444A1 GENERAL INFORMATION: APPLICANT: Myriad Genetics, Inc. APPLICANT: Zavitz, Kenton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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3 RPFAAAKE 10
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Best Local Similarity
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LENGTH: 10
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FEATURE:
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US-10-601-837-103

i Sequence 103, Application US/10601837

j Publication No. US20040053309A1

j GENERAL INFORMATION:
    APPLICANT: Holt, Gordon D

j APPLICANT: Kelly, Michael D

j APPLICANT: Kennedy, Sandra J

j TITLE OF INVENTION: Proteins, Genes and Their Use for Diagnosis and Treatment of Kidn

j TITLE OF INVENTION: Response

j TILE REFERENCE: 2543-1-030

j CURRENT APPLICATION NUMBER: US/10/601,837

j CURRENT APPLICATION NUMBER: US/0/2603/05777

j PRIOR APPLICATION NUMBER: US 60/260392

j PRIOR APPLICATION NUMBER: US 60/260392

j PRIOR APPLICATION NUMBER: US 60/260392

j NUMBER OF SEQ ID NOS: 272

j NUMBER OF SEQ ID NOS: 272

j NUMBER OF SEQ ID NOS: 272
                                                                                                                                                                                              Sequence 9, Application US/0995388

Fublication No. US20020106325A1

GENERAL INFORMATION:

APPLICANT: Carpenter, Jr., Alan P.

TITLE OF INVENTION:

TITLE OF INVENTION: RECEPTOR TARGETED IMAGING AGENT

FILE REFERENCE: BMS-2201

CURRENT APPLICATION NUMBER: US/09/995,388

CURRENT APPLICATION NUMBER: US 60/253,324

PRIOR PILING DATE: 2000-11-27

NUMBER OF SEQ ID NOS: 49

SOFTWARE: Patentin version 3.2

SEQ ID NO 9

LENGTH: 11
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Mismatches
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ORGANISM: Artificial Sequence
6; Conservative
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5 GDYVRGD 11
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LENGTH: 12
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Matches
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RESULT 42
UG-10-369-405-19
i Sequence 19, Application US/10369405
publication No. US20030224400A1
i GENERAL INFORMATION:
    APPLICANT: Bristol-Myers Squibb Company
    TITLE OF INVENTION: NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR, HGPREMY11, AND VARIANTS
    TITLE OF INVENTION: THEREOF
    PILE REFERENCE: D0075A CIP
    CURRENT APPLICATION NUMBER: US/10/369,405
    CURRENT APPLICATION NUMBER: US. 60/249,613
    PRIOR RELING DATE: 2000-11-17
    PRIOR PILING DATE: 2000-11-17
    PRIOR FILING DATE: 2001-11-16
    PRIOR FILING DATE: 2001-11-16
    PRIOR FILING DATE: 2001-12-21
    PRIOR FILING DATE: 2001-12-21
    PRIOR FILING DATE: 2001-12-21
    PRIOR FILING DATE: 2001-10-16
    PRIOR FILING DATE: 2001-10-16
    PRIOR FILING DATE: 2001-10-16
    PRIOR FILING DATE: 2001-10-16
    PRIOR FILING DATE: 2001-07-16
    NUMBER OF SEQ 1D NOS: 94
    SOFTWARE: PatentIn version 3.2
    SEQ ID NO 19
TITLE OF INVENTION: A NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR, HGPRBMY11, EXPRESSED HIC TITLE OF INVENTION: HEART AND VARIANTS THEREOF FILE REFERENCE: D0075.NP CURRENT APPLICATION NUMBER: US/09/991,225 CURRENT PILING DATE: 2001-11-16 PRIOR PILING DATE: 2001-11-17 PRIOR FILING DATE: 2000-11-17 PRIOR FILING DATE: 2000-11-27 PRIOR FILING DATE: 2000-12-21 PRIOR FILING DATE: 2000-12-21 PRIOR FILING DATE: 2001-07-16 NUMBER: 60/305,818 PRIOR FILING DATE: 2001-07-16 NUMBER OF SEQ ID NOS: 81 SOFTWARE: Patentin version 3.0 SEQ ID NO 38 LENGTH: 13
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Publication No. US20030224400A1

GENERAL INFORMATION:
APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR, HGPRBMY11, AND VARIANTS
TITLE OF INVENTION: THEREOF
FILE REFERENCE: D0075A CIP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 13;
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Pred. No. 1.2e+04;
6; Mismatches 0; Indels
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Best Local Similarity 33.5
Local 3; Conservative
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Best Local Similarity 33.3
Matches 3; Conservative
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CRCANISM: homo sapiens
US-10-369-405-19
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4 VTSIRSAWI 12
                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT ORGANISM: Homo sapiens
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US-10-369-405-38
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; Sequence 19, Application US/09991225
; Publication No. US20030153063A1
; GENERAL INFORMATION:
APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: HEART AND VARIANTS THEREOF
; TITLE OF INVENTION: HEART AND VARIANTS THEREOF
; FILE PEPERROCE: D0075.NP
; CURRENT APPLICATION NUMBER: US/09/991,225
; CURRENT FILING DATE: 2001-11-16
; PRIOR FILING DATE: 2000-11-17
; PRIOR FILING DATE: 2000-11-17
; PRIOR FILING DATE: 2000-12-21
; PRIOR FILING DATE: 2001-07-16
; NUMBER OF SEQ ID NOS: 81
; SOFTWARE: PatentIn version 3.0
; FEACTOR 19
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                                                                        Sequence 2922, Application US/10224999A;
Sequence 2922, Application US/10224999A;
Publication No. US20030171318A1;
GENERAL INFORMATION:
APPLICANT: Morham, Scott
APPLICANT: Morham, Scott
APPLICANT: Application and Method for Treating Viral Infection
TITLE OF INVENTION: Composition and Method for Treating Viral Infection
FILE REFERENCE: 5004.01
CURRENT FILING DATE: 2003-03-03
CURRENT FILING DATE: 2003-03-03
CURRENT FILING DATE: 2001-08-20
NUMBER OF SEQ ID NOS: 3484
SOFTWARE: Patentin version 3.1
SEQ ID NO 2922
LENGTH: 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 32; DB 10; Length 13
Pred. No. 1.2e+04;
5; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            33.3%;
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Best Local Similarity 33.3
Matches 3; Conservative
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4 VTSIRSAWI 12
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CRGANISM: homo sapiens
US-09-991-225-19
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Matches 6; Conserv
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Sequence 2931, Application US/10224999A

Publication No. US20030171318A1

GENERAL INFORMATION:

APPLICANT: Morham, Scott

APPLICANT: Zavitz, Kenton

APPLICANT: Eavitz, Kenton

TITLE OF INVENTION: Composition and Method for Treating Viral Infection

FILE REFERENCE: 5004.01

CURRENT APPLICATION NUMBER: US/10/224,999A

CURRENT FILING DATE: 2003-03-03

PRIOR FILING DATE: 2001-08-20

NUMBER OF SEQ ID NOS: 3484

SOFTWARE: Patentin version 3.1

SEQ ID NO 2931

LENGTH: 13
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Sequence 2932, Application US/10224999A

Fublication No. US20030171318A1

GENERAL INFORMATION:

APPLICANT: Myriad Genetics, Inc.

APPLICANT: Apolication Scott

APPLICANT: Apolication and Method for Treating Viral Infection

TITLE OF INVENTION: Composition and Method for Treating Viral Infection

FILE REFERENCE: 5004.01

CURRENT APPLICATION NUMBER: US/10/224,999A

CURRENT FILING DATE: 2003-03

PRIOR APPLICATION NUMBER: US 60/313,695
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Pred. No. 1.2e+04;
1; Mismatches 5; Indels
; CURRENT APPLICATION NUMBER: US/10/369,405
; CURRENT FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: U.S. 60/249,613
; PRIOR FILING DATE: 2000-11-17
; PRIOR FILING DATE: 2001-11-16
; PRIOR FILING DATE: 2001-11-16
; PRIOR FILING DATE: 2001-12-21
; PRIOR FILING DATE: 2000-12-21
; PRIOR FILING DATE: 2001-07-16
; NUMBER OF SEQ ID NOS: 94
; SOFTWARE: PATENTIN Version 3.2
; SEQ ID NO 38
; LENGTH: 13
; TYPE: PRT

; ORGANISM: Homo sapiens
US-10-369-405-38
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Best Local Similarity 50.0%;
Matches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 3.0
Best Local Similarity 33.3
Matches 3; Conservative
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4 VTSIRSAWI 12
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US-10-224-999A-2932
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Best Local Similarity 87.5
Matches 7; Conservative
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USA
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STATE: MA
COUNTRY: US/
ZIP: 02110
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Query Match
8
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Sequence 10, Appl
Sequence 40, Appl
Sequence 120, Appl
Sequence 2, Appli
Sequence 101, Appl
Sequence 25, Appl
Sequence 25, Appl
Sequence 25, Appl
Sequence 23, Appl
Sequence 24, Appl
Sequence 36, Appl
Sequence 36, Appl
Sequence 36, Appl
Sequence 37, Appl
Sequence 36, Appl
Sequence 36, Appl
Sequence 8, Appl
Sequence 8, Appl
Sequence 8, Appl
                                                                                                                                                          August 30, 2004, 10:40:59; Search time 82.2162 Seconds (without alignments) 130.609 Million cell updates/sec
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1 MKVLLAAALIAGSVFFLLLP......VIIADCGKIEVEKPFAIAKE 208
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1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
                      GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-08-944-604-10
US-08-556-597-40
PCT-US93-06751-120
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US-07-963-324-2
US-07-963-324-2
US-08-448-418-101
US-08-146-979-101
US-08-973-559-25
US-08-973-235-5
US-08-993-235-5
US-08-993-235-5
US-08-475-041-23
US-08-475-041-23
US-08-475-041-23
US-08-475-041-23
US-08-361-864-27
US-08-361-864-27
US-08-361-864-27
US-08-361-864-27
US-08-111-127-19
US-08-658-639-8
US-08-658-639-8
US-08-658-639-8
US-08-658-606-26
US-08-973-563A-27
US-08-973-563A-27
US-08-973-563A-27
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Maximum Match 100%
Listing first 45 summaries
                                                                                                               - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 14
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APPLICANT: KEESEE, SUSAN
APPLICANT: OBAR, ROBERT
APPLICANT: WU, YING-JYE
TITLE OF INVENTION: MATERIALS AND METHODS FOR DETECTION OF
TITLE OF INVENTION: BREAST CANCER
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Testa, Hurwitz & Thibeault
STREET: 125 High St.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.5%; Score 38; DB 2; Length 9; 87.5%; Pred. No. 3e+05;
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/658,639
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: MEYERS, THOMAS C
REGISTRATION NUMBER: 36,989
REGISTRATION NUMBER: 36,989
REFERENCE/DOCKET NUMBER: MTP-021 (8395/24)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
TELEFAX: (617) 248-7100
JINFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
TENNITH: 9 amino acids
US-08-993-235-7

US-08-993-235-7

US-08-916-913A-4

US-09-591-564-7

US-09-591-564-7

US-09-591-564-7

US-08-271-830-56

US-08-271-830-56

US-08-416-870C-15

US-08-466-151-50

US-08-466-151-50

US-08-466-151-17

US-08-466-151-17

US-08-466-151-46

US-08-466-151-46

US-08-466-151-46

US-08-466-153B-17

US-08-466-153B-17

US-08-466-153B-17

US-08-466-153B-17

US-08-466-153B-17

US-08-466-153B-17

US-08-466-153B-17
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                                      Sequence 10, Application US/08944604

Patent No. 6218131

GENERAL INFORMATION:
APPLICANT: KEESEE, SUSAN
APPLICANT: OBAR, ROBERT
APPLICANT: OBAR, ROBERT
TITLE OF INVENTION: MATERIALS AND METHODS FOR DETECTION OF
TITLE OF INVENTION: BREAST CANCER
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: Testa, Hurwitz & Thibeault
STREET: 125 High St.
CITY: Boston
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ·.
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Fatent No. 5877155

GENERAL INFORMATION:

APPLICANT: Miller, Jonathan L.

APPLICANT: Lyle, Vicki A.

ITILE OF INVENTION: MIMOTOPES AND ANTI-MIMOTOPES OF

TITLE OF INVENTION: HUMAN PLATELET GLYCOPROTEIN Ib/IX

NUMBER OF SEQUENCES: 173

CORRESPONDENCE ADDRESS:

ADDRESSEE: Nixon, Hargrave, Devans & Doyle LLP

STREET: Clinton Square, P.O. Box 1051

CITY: Rochester

STATE: New York

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 38; DB 3; Length 9; Pred. No. 3e+05;
                                                                                                                                                                                                                                                                                                                                                                          ZIP: 0210

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/944,604

FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MTP-021 (8395/24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MEYERS, THOMAS C
REGISTRATION NUMBER: 36,989
REFERENCE/DOCKET NUMBER: MTP-O
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 3.5%;
Best Local Similarity 87.5%;
Matches 7; Conservative
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                189 VIIADCGK 196
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RESULT 2
US-08-944-604-10
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                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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Sequence 120, Application PC/TUS9306751
GENERAL INFORMATION:
APPLICANT: P. Keller, A. J. Conley, A.R. Shaw, B.A. Arnold
TITLE OF INVENTION: Immunological Conjugates of OMPC and
TITLE OF INVENTION: HIV-Specific Selected Principal Neutralization GXG Epitopes
NUMBER OF SEQUENCES: 146
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READBLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Fatentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/06751
FILING DATE: 19930719
CLASSIFICATION:
NAME: Meredith, Roy D.
REGISTRATION NUMBER: 30,777
REFERENCE/DCCKET NUMBER: 18614
TELEPHONE: (908) 594-4678
TELEPHONE: (908) 594-4720
metrov 13003
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/556,597
                                                                                             APFLICATION: 530

PRICING DATE:
CLASSIFICATION: 530

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/406,330
FILLING DATE: 17-MAR-1995
ATTORNEY AGENT INFORMATION:
NAME: Timian, Susan J.
REGISTRATION NUMBER: 34,103
REFERENCE/DOCKET NUMBER: 20884/101
TELEPHONE: (716) 263-1636
TELEPHONE: (716) 263-1636
TELEFAX: (716) 263-1636
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARRACTERISTICS:
LENGTH: 12 amino acids
""""" amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Merck & Co., Inc. STREET: P.O. Box 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (908) 594-4720
TELEX: 138825
INFORMATION FOR SEQ ID NO: 120:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 3.4%;
Best Local Similarity 66.7%;
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                156 GKHVVFGKVLEG 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: sin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: P.O. CITY: Rahway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: NJ
COUNTRY: US
ZIP: 07065
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Holliger, Kaspar-Philipp
Griffiths, Andrew D
Hoogenboom, Hendricus RJM
Malmqvist, Magnus
Marks, James D
McGuinness, Brian T
Pope, Anthony R
NAME: Bencen, Gerard H
REGISTRATION NUMBER: 35,746
REFERENCE/DOCKET NUMBER: 18787
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908)594-3901
TELEFAX: (908)594-4720
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 101, Application US/08448418; Patent No. 5837242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Prospero, Terence D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 85.7
Matches 6; Conservative
                                                                                                                                                                                                                                                                                           FRAGMENT TYPE: internal FEATURE:
                                                                                                                                                                  LENGTH: 9 amino acids TYPE: AMINO ACID
                                                                                                                                                                                                         TOPOLOGY: both
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 122 LKHYGPG 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 LKHIGPG 7
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
APPLICANT:
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Patent No. 6255055
GENERAL INFORMATION:
APPLICANT: Ross, Jeffrey
TITLE OF INVENTION: (CRD-BP) AND ITS NUCLEIC ACID SEQUENCE
FILE REFERENCE: 960296.95131
CURRENT APPLICATION NUMBER: US/09/261,855A
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 46
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 9
LENGTH: 13
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Sequence 2, Application US/07963324;

Patent No. 5274122;

GENERAL INFORMATION:

APPLICANT: Marburg, Stephen

APPLICANT: Leanza, William J

TITLE OF INVENTION: Acidic Derivatives of Homocysteine

TITLE OF INVENTION: Thiolactone

NUMBER OF SEQUENCES: 4

CORRESPONDENCE ADDRESS:

ADDRESSEE: Merck & Co., Inc.

STREET: 126 Lincoln Avenue

CITY: Rahway

STATE: New Jersey

COUNTRY: United States of America

ZIP: 07065
                                                                                                                          Query Match 3.4%; Score 37; DB 5; Length 14; Best Local Similarity 55.6%; Pred. No. 1.1e+03; Matches 5; Conservative 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.3%; Score 36; DB 3; Length 13; 63.6%; Pred. No. 1.38+03; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Consensus sequence for SEQ ID NOs:3-8.
US-09-261-855-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READBLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/963,324
FILING DATE: 19921015
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
                                                        ; IMMEDIATE SOURCE: Random Epitope Library Beta PCT-US93-06751-120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 3.3
Best Local Similarity 63.6
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97 GDFTRGDGTGG 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 GGFGRGGGRGG 13
     MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                              120 FKLKHYGPG 128
                                                                                                                                                                                                                                        :: ||||||
3 YRAAHYGPG 11
                          HYPOTHETICAL:
ANTI-SENSE: N
                                                                                                                                                                                                                                                                                                                  RESULT 5
US-09-261-855-9
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Gaps
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; OTHER INFORMATION: /label= cycle
; OTHER INFORMATION: /note= "amide bond through Lys epsilon amino"
US-07-963-324-2
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TITLE OF INVENTION: Multivalent and Multispecific Binding
TITLE OF INVENTION: Proteins, Their Manufacture and Use
NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (BPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/48,418
FILING DATE: 14-MAY-1996
CLASSIFICATION: C12N 15/62, 15/70, C07K 1/00
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/GB93/02492
FILING DATE: 03-DEC-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                         1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Marshall O'Toole Gerstein Murray and Borun STREET: 6300 Sears Tower 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                  3.2%; Score 35; DB 1; Length 9;
85.7%; Pred. No. 3e+05;
tive 0; Mismatches 1; Indels
NAME/KEY: Modified-site
LOCATION: 1
OTHER INFORMATION: /label= Nle
OTHER INFORMATION: /note= "norleucine"
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amino acid
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APPLICANT: Prospero, Terence D
APPLICANT: Winter, Gregory P
TITLE OF INVENTION: Multivalent and Multispecific Binding
TITLE OF INVENTION: Proteins, Their Manufacture and Use
NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Chicago
STATE: 11linois
COUNTRY: USA
ZIP: 60606-6402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/146,979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 3.2%; Score 35; DB 2; Length 14; Best Local Similarity 50.0%; Pred. No. 1.8e+03; Matches 7; Conservative 2; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 14;
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PRILICATION NUMBER: GB 9225453.1
FILING DATE: 04-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9300816.7
FILING DATE: 16-JAN-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 93303614.7
FILING DATE: 10-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9319969.3
FILING DATE: 12-SEP-1993
ATTORNEY/AGENT INFORMATION:
NAME: DAVIG W. Clough
REGISTRATION NUMBER: 36,107
REFERENCE/DOCKET NUMBER: 28111/32651
INFORMATION FOR SEQ ID NO: 101:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 aming acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/448,418
FILING DATE: 14-MAY-1996
APPLICATION NUMBER: PCT/GB93/02492
FILING DATE: 03-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9225453.1
FILING DATE: 04-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Holliger, Kaspar-Philipp
APPLICANT: Griffiths, Andrew D
APPLICANT: Hoogenboom, Hendricus RJM
APPLICANT: Malmqvist, Magnus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 101, Application US/09146979 Patent No. 6492123 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Marks, James D
McGuinness, Brian T
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: peptide linker
US-08-448-418-101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              96 GGDFTRGDGTGGKS 109
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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Patent No. 5885965
GENERAL INFORMATION:
APPLICANT: Oppenheim, Frank G.
APPLICANT: Xu, Tao
APPLICANT: Spacciapoli, Peter
APPLICANT: Roberts, F. D.
APPLICANT: Friden, Philip M.
TITLE OF INVENTION: Anti-Fungal D-Amino Acid Histatin-Based
TITLE OF INVENTION: Peptides
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 35; DB 4; Length 14; Pred. No. 1.8e+03; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,563A
FILING DATE: 07-JUN-1996
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/485,273
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-02A2
TELEPHONE: 781-861-6240
TELEFAX: 781-861-6240
TELEFAX: 781-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
PRIOR APPLICATION DATA:

APPLICATION NUMBER: GB 9300816.7

FILING DATE: 16-JAN-1993

PRIOR APPLICATION DATA:

APPLICATION NUMBER: EP 93303614.7

FILING DATE: 10-MAY-1993

PRIOR APPLICATION DATA:

APPLICATION NUMBER: GB 9319969.3

FILING DATE: 22-SEP-1993

ATTORNEY/AGENT INFORMATION:

NAME: David W. Clough

REGISTRATION NUMBER: 36,107

REFERENCE/DOCKET NUMBER: 28111/32651

INFORMATION FOR SEQ ID NO: 101:

SEQUENCE CHARACTERISTICS:

LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear

MOLECULE TYPE: peptide linker
US-09-146-979-101
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INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 50.09
Matches 7, Conservative
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APPLICANT: OPPENHEIM, FRANK G. APPLICANT: ROBERTS, F. DONALD
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Best Local Similarity 58.3
Matches 7; Conservative
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Matches 7; Conservative
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2 KRHFGYKR-KFH 12
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2 KRHFGYKR-KFH 12
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             2 KRHFGYKR-KFH 12
                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
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                                                                                                                                                                                                    3.1%; Score 33.5; DB 2; Length 12; 58.3%; Pred. No. 2.1e+03; iive 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 25, Application US/08973559;
Patent No. 5912230;
GENERAL INFORMATION:
APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: KOBERTS, F. D.
APPLICANT: SPACCIAPOLI, PETER
APPLICANT: FRIDEN, PHILIP M.
TITLE OF INVENTION: Histatin-Based Peptides
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
STREET: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
                                                                                                         LOCATION: 1..12;
COTHER INFORMATION: /note= "At least one amino acid;
COTHER INFORMATION: must have a D configuration."
US-08-973-563A-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,559
FILING DATE: 07-JUN-1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/481,888
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PER95-01A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 781-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 781-861-6240
TELEFAX: 781-861-9540
INFORMATION FOR SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 3.1
Best Local Similarity 58.3
Matches 7; Conservative
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Matches 7; Conservative
                                                                                                                                                                                                                                                                               75 EKGFGYKNSKFH 86
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2 KRHFGYKR-KFH 12
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                                TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 02173
COMPUTER READABLE FORM:
STRANDEDNESS:
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                                                                                          NAME/KEY: Region
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US-08-973-559-25
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75 EKGFGYKNSKFH 86

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101 RGDGTGG 107
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Matches 6; Conserv
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US-08-475-041-23
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Sequence 23, Application US/08273274
Sequence 5849260
GENERAL INFORMATION:
APPLICANT: Dean, Richard T
APPLICANT: Lister-James, John
TITLE OF INVENTION: Technetium-99m Labeled Peptides for TITLE OF INVENTION: Thrombus Imaging
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.0%; Score 33; DB 1; Length 8; 85.7%; Pred. No. 3e+05; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,880
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
ATTONNEY/AGENT INFORMATION:
NAME: No. 5736122nan, Kevin E
REGISTRATION NUMBER: 92,216-L
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-715-1234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSE: Allegretti & Witcoff, Ltd.
STREET: 10 South Wacker Drive Suite 3000
CITY: Chicago
STATE: 11linois
CONTRY: USA
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/273,274
                  ADDRESSEE: Banner & Allegretti, Ltd.
STREET: 10 South Wacker Drive Suite 3000
CITY: Chicago
STATE: Illinois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 436
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/886,752
FILING DATE: 21-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: No. 5849260nan, Kevin E
REGISTRATION NUMBER: 35,303
                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 312-715-1234
TELEX: 910-221-5317
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 3.0
Best Local Similarity 85.7
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear;
MOLECULE TYPE: peptide
US-08-482-880-23
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              101 RGDGTGG 107
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GY: linear
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                                                                                                                    90909
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 23, Application US/08475041
Patent No. 5879658
GENERAL INFORMATION:
APPLICANT: Dean, Richard T
APPLICANT: Lister-James, John
TITLE OF INVENTION: Technetium-99m Labeled Peptides for TITLE OF INVENTION: Thrombus Imaging
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.0%; Score 33; DB 2; Length 8; 85.7%; Pred. No. 3e+05; 1; Indels ive
                                                                                                                                                                                                                                                                                    Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Perentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/475,041
FILING DATE: 07-UN-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: No. 5879658nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 92,216-M
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-715-1000
                                                                                                                                                                                                                                                                                 Score 33; DB 2;
Pred. No. 3e+05;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: Banner & Allegretti, Ltd.
STREET: 10 South Wacker Drive Suite 3000
CITY: Chicago
STATE: 111inois
COUNTRY: USA
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-715-1000
TELEFAX: 312-715-1234
TELEX: 910-221-5317
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                 Query Match 3.0%;
Best Local Similarity 85.7%;
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 312-715-1234
TELEX: 910-221-5317
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                  TYPE: amino acids to TOPOLOGY: linear i MOLECULE TYPE: peptide US-08-273-274-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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MOLECULE TYPE: peptide
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Sequence 23, Application US/08335832

Factor No. 5925331

GENERAL INFORMATION:

APPLICANT: Dean, Richard T

APPLICANT: Lister-James, John

TITLE OF INVENTION: Technetium-99m Labeled Peptides for TITLE OF INVENTION: Thrombus Imaging

NUMBER OF SEQUENCES: 53

CORRESPONDENCE ADDRESS:

ADDRESSEE: Banner & Allegretti, Ltd.

STREET: 10 South Wacker Drive Suite 3000

CITY: Chicago

STATE: Illinois

COUNTRY: USA

ADDRESSE: CORRESPONDENCE DRIVE SUITE 3000

CITY: Chicago

STATE: Illinois

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.0%; Score 33; DB 2; Length 8; 85.7%; Pred. No. 3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER CEADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/335,832
FILING DATE: 05-JAN-1995
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/361,864
FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/955,466A
FILING DATE: 19921002
ATTORNEY/AGENT INFORMATION:
NAME: NO. 5977064nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 92,668
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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NAME: No. 5925331nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 92,216-I
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-715-1000
TELEX: 312-715-1234
TELEX: 310-221-5317
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                         TELEX: 910-221-5317
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
GY: linear
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Best Local Similarity
Matches 6; Conserv
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TOPOLOGY:
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| Sequence 27, Application US/08361864
| Patent No. 5977064
| Patent No. 5977064
| APPLICANT: Dean, Richard T APPLICANT: Lister-James, John TITLE OF INVENTION: Multimeric Polyvalent Antithrombotic TITLE OF INVENTION: Agents NUMBER OF SEQUENCES: 37
| CORRESPONDENCE ADDRESS: ADDRESSE: Allegretti & Witcoff, Ltd. STREET: 10 South Wacker Drive, Suite 3000 CITY: Chicago STATE: Illinois
| STATE: Illinois | STATE: USA
                                                                                                GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Dean, Richard T
APPLICANT: Lister-James, John
TITLE OF INVENTION: Technetium-99m Labeled Peptides for
TITLE OF INVENTION: Thrombus Imaging
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Allegretti, Ltd.
STREET: 10 South Wacker Drive Suite 3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.0%; Score 33; DB 2; Length 8; 85.7%; Pred. No. 3e+05; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT SISIEM: FC-DOS/MS-LOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,773
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: No. 5968476nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 92,216-O
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRAX: 312-715-1234
TELEX: 910-221-5317
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC_compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk

...ntmeR: IBM PC compatible
...ntmeR: TPM PC compatible
                               US-08-484-773-23; Sequence 23, Application US/08484773; Patent No. 5968476
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Best Local Similarity 85.7
Matches 6; Conservative
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MOLECULE TYPE: peptide
US-08-484-773-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      101 RGDGTGG 107
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                                                                                                                                                                                                                                                                                                                       CITY: Chicago
STATE: Illinois
COUNTRY: USA
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Patent No. 5951981
GENERAL INFORMATION:
APPLICANT: Markland Jr., Francis S.
APPLICANT: Bush, Larry R.
APPLICANT: Swenson, Stephen
APPLICANT: Flores Sanchez, Eladio
TITLE OF INVENTION: THROMBOLYTIC AGENTS WITH ANTITHROMBOTIC ACTIVITY
FILE REFERENCE: DIT1 124
CURRENT APPLICATION NUMBER: US/08/753,781C
CURRENT FILING DATE: 1996-12-02
NUMBER OF SEQ ID NOS: 40
SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTHER INFORMATION: Description of Artificial Sequence:synthetic; CTHER INFORMATION: peptide US-09-141-127-19
                                        1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Indels
 DB 2; Length 9; 3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 33; DB 2; Length 9;
Pred. No. 3e+05;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 9;
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Pred. No. 3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INC. SUGGATION:

GENERAL INC. SUGGATION:

APPLICANT: Dean, Richard T.

APPLICANT: Lister-James, John

TITLE OF INVENTION: THROMBUS IMAGING AGENTS;
FILE REFERENCE: DITI 113.1USC1

CURRENT APPLICATION NUMBER: US/09/141,127A;
CURRENT FILING DATE: 1998-08-27

EARLIER APPLICATION NUMBER: PCT/US93/04794;
EARLIER FILING DATE: 1995-01-05;
EARLIER PILING DATE: 1993-05-21

BARLIER APPLICATION NUMBER: 07/886,752

EARLIER FILING DATE: 1993-05-21

SARLIER FILING DATE: 1993-05-21

HOWBER OF SEQ ID NOS: 40

SOFTWARE: Patentin Ver. 2.0
                                        0; Mismatches
   Score 33;
Pred. No.
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 3.0%;
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                        6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6; Conservative
                                                                               101 RGDGTGG 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           101 RGDGTGG 107
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Best Local Similarity
Query Match
Best Local Similarity
Matches 6; Conserv
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Best Local Similarity
Matches 6; Conserv
                                                                                                                    2 RGDGGGG 8
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US-08-753-781-36
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US-09-141-127-19
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LENGTH: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
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US-08-944-604-8
; Sequence 8, Application US/08944604
; Patent No. 6218131
; GENERAL INFORMATION:
; APPLICANT: KEESEE, SUSAN
; APPLICANT: OBAR, ROBERT
; TITLE OF INVENTION: MATERIALS AND METHODS FOR DETECTION OF
; TITLE OF INVENTION: BREAST CANCER
; NUMBER OF SEQUENCES: 24
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Testa, Hurwitz & Thibeault
; STREET: 125 High St.
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                                                                                                                     RESULT 21
US-08-658-639-8
; Sequence 8, Application US/08658639
; Patent No. 5914238
; GENERAL INFORMATION:
APPLICANT: KEESEE, SUSAN
; APPLICANT: OBERT
; TITLE OF INVENTION: MATERIALS AND METHODS FOR DETECTION OF
; TITLE OF INVENTION: BREAST CANCER
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 33; DB 2; Length 10; Pred. No. 1.9e+03; 2; Mismatches 2; Indels
 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/658,639
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: MEYERS, THOMAS C
REGISTRATION NUMBER: 36,989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 36,989
REFERENCE/DOCKET NUMBER: MTP-021 (8395/24)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Mismatches
                                                                                                                                                                                                                                                                                                                                        E: Testa, Hurwitz & Thibeault
125 High St.
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 10 amino acids
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                    101 RGDGTGG 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
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Matches 6; Conserv
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STREET: 12.
"V' Boston
                                                                     2 RGDGGGG
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Matches
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us-09-720-469a-44.aug30.rai

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Query Match
Best Local Similarity 100.
Thes 6; Conservative
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CITY: San Francisco
STATE: California
COUNTRY: United State
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: not
                                                                                                                                                                                                                                                                                                                                                   28 KKKGPK 33
                                                                                                                                                                                                                                                                                                                                                                                        1 KKKGPK 6
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US-08-656-906-26
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TITLE OF INVENTION: Inhibition of HIV Replication Using Soluble Tat Peptide Analogs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.0%; Score 33; DB 3; Length 10; 60.0%; Pred. No. 1.9e+03; iive 2; Mismatches 2; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/269,991B
FILING DATE: 07-Sep-1999
CLASSIFICATION: <UNKNOWn>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/944,604
                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: MEYERS, THOMAS C
REGISTRATION NUMBER: 36,989
REFERENCE/DOCKET NUMBER: MTP-021 (8395/24)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/17704
FILLING DATE: 02-OCT-1997
APPLICATION NUMBER: US 60/027,658
FILING DATE: 04-OCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: Garrett-Wackowski, Eugenia
REGISTRATION NUMBER: 37,330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Kashanchi, Fatah
Sadaie, Mohamad Reza
Brady, John N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 10, Application US/09269991B Patent No. 6686333 GENERAL INFORMATION:
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ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   114 RFPDENFKLK 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         : | | | | | | :
1 KFDDENFILR 10
                                                                                                                                                                                                             FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 6; Conserv
           MA
USA
Boston
                                                       02110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-269-991B-10
               STATE: M2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-944-604-8
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/656,906
FILING DATE: 03-JUN-1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 23-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO WO 95/25809
FILING DATE: 23-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/216,534
FILING DATE: 23-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Ingolia, Diane E.
REGISTRATION NUMBER: 40,027
REFERENCE/DOCKET NUMBER: 40,027
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRAX: (415) 397-8338
INFORMATION FOR SEQ ID NO: 26:
REFERENCE/DOCKET NUMBER: 015280-276100US
                                                                                                                              TYPE: amino acid
STRANDEDNESS: «Unknown»

TOPOLOGY: linear

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION: SEQ ID NO: 10:
US-09-269-991B-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    United States Of America
              TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 aming acids
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TELEFAX: 781-861-9540
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
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Best Local Similarity 77.00
1.20 7; Conservative
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OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
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 US-08-973-563A-27
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US-08-973-559-27
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                                                                                Gaps
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                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ·.
                                      Query Match 3.0%; Score 33; DB 2; Length 14; Best Local Similarity 50.0%; Pred. No. 2.9e+03; Matches 7; Conservative 2; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.0%; Score 33; DB 3; Length 14; ilarity 50.0%; Pred. No. 2.9e+03; Conservative 2; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                               APPLICANT: Ferkol Jr., Thomas W.
APPLICANT: Ferkol Jr., Thomas W.
APPLICANT: Davis, Pamela B.
APPLICANT: Ziady, Assem-Galal
TITLE OF INVENTION: Serpin Enzyme Complex Receptor
TITLE OF INVENTION: Mediated Gene Transfer
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: Medlen & Carroll
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/217,847
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/656,906
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Medlen & Carroll
STREET: 220 Montgomery Street, Suite 2200
CITY: San Francisco
STATE: California
COUNTRY: United States Of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION DATA:
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO WO 95/25809
FILING DATE: 23-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/216,534
FILING DATE: 23-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Ingolia, Diane E.
REGISTRATION NUMBER: 40,027
REFERENCE/DOCKET NUMBER: CASE-02280
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 397-8338
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
TENGRALION FOR SEQ ID NO: 26:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                        US-09-217-847-26; Sequence 26, Application US/09217847; Patent No. 6200801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SS: not relevant not relevant
                                                                                                                     96 GGDFTRGDGTGGKS 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96 GGDFTRGDGTGGKS 109
                                                                                                                                                          1 GGGGSGGGGGGS 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: peptide US-09-217-847-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
US-08-656-906-26
                                                                                                                                                         d
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Gaps
Sequence 27, Application US/08973563A

Sequence 27, Application US/08973563A

Patent No. 5885965

GENERAL INFORMATION:

APPLICANT: Oppenheim, Frank G.

APPLICANT: Spacciapoli, Peter

APPLICANT: Roberts, F. D.

APPLICANT: Friden, Philip M.

TITLE OF INVENTION: Anti-Fungal D-Amino Acid Histatin-Based

TITLE OF INVENTION: Peptides

NUMBER OF SEQUENCES: 37

CORRESPONDENCE ADDRESS:

ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.

STREET: Two Militia Drive

CORRESPONDENCE ADDRESS:

ADDRESSEE: Lexington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "At least one amino acid
must have a D configuration."
                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: US

COUNTRY: US

ZIP: 02173

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/973,563A

FILING DATE: 07-JUN-1996

CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 27, Application US/08973559
Patent No. 5912230
GENERAL INFORMATION:
APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: ROBERTS, F. D.
APPLICANT: SPACCIAPOLI, PETER
APPLICANT: FRIDEN, PHILIP M.
TITLE OF INVENTION: Anti-Fungal and Anti-Bacterial
TITLE OF INVENTION: Histatin-Based Peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 32.5; DB 2;
Pred. No. 2.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/485,273
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-02A2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 781-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PER95-02A2
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RESULT 26

us-09-720-469a-44.aug30.rai

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Best Local Similarity 77.8
Matches 7; Conservative
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ORGANISM: Homo sapiens
US-08-993-235-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 FGYKR-KFH 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              55402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: MC
COUNTRY:
ZIP: 554
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-394-748A-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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US-08-993-235-7
Sequence 7, Application US/08993235
Patent No. 6084064
GENERAL INFORMATION:
APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: ROBERTS, F. DONALD
APPLICANT: SPACCIAPOLI, PETER
TILLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES
FILE REFERENCE: 50032/002001
CURRENT FILING DATE: 1997-12-18
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.0%; Score 32.5; DB 3; Length 12; 77.8%; Pred. No. 2.7e+03; Live 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.0%; Score 32.5; DB 2; Length 12; 77.8%; Pred. No. 2.7e+03; tive 0; Mismatches 1; Indels
                                                                                                                                         ZIP: 02173

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,559
FILING DATE: 07-JUN-1996
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/481,888
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-01A2
REGISTRATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 781-861-6240
NUMBER OF SEQUENCES: 37

CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
STATE: MA
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 781-861-9540
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 77.8
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Homo sapiens
US-08-993-235-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               78 FGYKNSKFH 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 FGYKR-KFH 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid STRANDEDNESS:
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Best Local Similarity
Matches 7; Conserv
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Sequence 4, Application US/08394748A

Fatent No. 6013628

GENERAL INFORMATION:

APPLICANT: Skubitz, Amy P.N.

APPLICANT: Furcht Leo T.

APPLICANT: Balles, Mark

APPLICANT: Agarwal, Anita

APPLICANT: Agarwal, Anita

APPLICANT: Murali, Shobana

TITLE OF INVENTION: Using Polypeptides

NUMBER OF SEQUENCES: 16

CORRESPONDENCE ADDRESS:

ADDRESSEE: Merchant & Gould
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.0%; Score 32.5; DB 4; Length 12; 77.8%; Pred. No. 2.7e+03; tive 0; Mismatches 1; Indels
                                                                      APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: OPPENHEIM, F. DONALD
APPLICANT: XU, TAO
APPLICANT: SPACCIAPOLI, PETER
TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES
FILE REFERENCE: 50032/002001
CURRENT APPLICATION NUMBER: US/08/993,235
CURRENT FILING DATE: 1997-12-18
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FASESEQ for Windows Version 3.0
SEQ ID NO 7
LENGTH: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/394,748A
FILING DATE: 27-FEB-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/203,458
FILING DATE: 28-FEB-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Carter, Charles G.
REGISTRATION NUMBER: 35,093
REFERENCE/DOCKET NUMBER: 600.307US01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-332-5300
TELEFAX: 612-332-9081
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Merchant & Gould
STREET: 3100 No. 6013628west Center
CITY: Minneapolis
Sequence 7, Application US/08993235
Patent No. 6531573
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:

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RESULT 33
PCT-US95-02478-4
                               RESULT 32
US-09-591-564-7
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Pred. No. 3e+03;
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Pred. No. 3e+03;
2; Mismatches 0; Indels
                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Fredrikson & Byron, P.A.
STREET: 900 Second Avenue South
CITY: Minneapolis
STATE: Minnesota
COUNTRY: USA
ZIP: 55402-3397
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage
COMPUTER: IBM Compatible
COMPUTER: ASCII files
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/916,913A
FILING DATE: 15 August 1997
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Clapper, David L.
APPLICANT: Swanson, Melvin J.
APPLICANT: Hu, Sheau-Ping
APPLICANT: Hu, Sheau-Ping
APPLICANT: Hos, Richard A.
APPLICANT: Everson, Terrence P.
TITLE OF INVENTION: LATENT REACTIVE POLYMERS WITH
TITLE OF INVENTION: BIOLOGICALLY ACTIVE MOIETIES
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                           .
                                                                                                                                                                                                          2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: No. 6121027e
FILING DATE: No. 6121027e
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                        Sequence 7, Application US/08916913A Patent No. 6121027
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13 amino acid residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Goldman, Philip M.
REGISTRATION NUMBER: 31,162
REFERENCE/DOCKET NUMBER: 9896.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (612) 347-7077
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.0%;
                                                                                                                                                                     Query Match 3.0%;
Best Local Similarity 70.0%;
Matches 7; Conservative
TYPE: amino acid
TOPOLOGY: 11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 70.0
Matches 7; Conservative
           SEQUENCE CHARACTERISTICS
                                                                            MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                            39 YFDLRI-GDE 47
                                                                                                                                                                                                                                                                                 4 YFDLRLKGDK 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         39 YFDLRI-GDE 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 YFDLRLKGDK 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION: APPLICANT: Clappe:
                                                                                                                   STRAIN: Hep-III
                                                                                                                                     US-08-394-748A-4
                                                                                                                                                                                                                                                                                                                                      RESULT 31
US-08-916-913A-7
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Gaps
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TITLE OF INVENTION: LATENT REACTIVE POLYMERS WITH BIOLOGICALLY ACTIVE MOIETIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1;
                                                                                                                                                                                                                                                                                                                                                             STATE: Minnesota
COUNTRY: USA
ZIP: 55402-3397
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows-97
SOFTWARE: ASCII files
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Method for Treating Conditions TITLE OF INVENTION: Of the Eye Using Polypeptides NUMBER OF SEQUENCES: 16
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.0%; Score 32.5; DB 4;
70.0%; Pred. No. 3e+03;
Live 2; Mismatches 0;
                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Fredrikson & Byron, P.A.
STREET: 900 Second Avenue South
CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/591,564
FILING DATE: 09-Jun-2000
CLASSIFICATION: <URNOwn>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/916,913
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 13 amino acid residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Goldman, Philip M.
REGISTRATION NUMBER: 31,162
REFERENCE/DOCKET NUMBER: 985
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4, Application PC/TUS9502478
GENERAL INFORMATION:
APPLICANT:
Sequence 7, Application US/09591564
Patent No. 6514734
GENERAL INFORMATION:
APPLICANT: Clapper, David L. Swanson, Melvin J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELBFAX: (612) 347-7077 INFORMATION FOR SEQ ID NO: 7: SEQUENCE CHARACTERISTICS:
                                                                                                                                                 Amos, Richard A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: single
                                                                                                                             Hu, Sheau-Ping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 70.0
Matches 7; Conservative
                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          39 YFDLRI-GDE 47
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Score 32; DB 1;
Pred. No. 3e+05;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                 Sequence 15, Application US/08416870C Patent No. 5824862 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 760-:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 205-8050
TELEFAX: (703) 205-8050
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MURPHY JR, GERALD M
REGISTRATION NUMBER: 28,97
  3.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 83.3
Matches 5; Conservative
        Query Match
Best Local Similarity 57.1
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
                                                                                                                                        126 GPGWVSM 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino acid
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                                                                                                                                                                                                  1 GPGWLDV 7
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                                                                                                                                                                                                                                                                                                                                        US-08-416-870C-15
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                                                                                                                                                                                                                                                                                                               RESULT 35
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US-08-271-830-56
; Sequence 56, Application US/08271830
; Patent No. 5510332
; GENERAL INFORMATION:
; APPLICANT: Kogan, Timothy P.
; APPLICANT: Ren, Kaijun
APPLICANT: Vanderslice, Peter
; APPLICANT: Beck, Pamela J.
; TITLE OF INVENTION: A PROCESS OF INHIBITING THE BINDING OF THE
; TITLE OF INVENTION: LINEAR PEPTIDES THEREFOR
; TITLE OF SEQUENCES: 102
; CORRESPONDENCE ADDRESS:
; CORRESPONDENCE ADDRESS:
; ANDMAR OF SEQUENCES: And A SEQUENCES:
; CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.0%; Score 32.5; DB 5; Length 13; 70.0%; Pred. No. 3e+03; tive 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Dressler, Goldsmith, Shore & Milnamow, Ltd. STREET: 180 No. 5510332th Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/271,830
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: /label= Xaa OTHER INFORMATION: /note= "Xaa=Pro-NH2." US-08-271-830-56
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FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: No. 5510332thrup, Thomas E.
REGISTRATION NUMBER: 33,268
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312)616-5400
TELEFAX: (312)
                APPLICATION NUMBER: PCT/US95/02478
                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/203,458
FILING DATE: 28-FEB-1994
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: Floppy disk
IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: Modified-site
                                                                                                                                                                                                                                                                            LENGTH: 13 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 70.0
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide ORIGINAL SOURCE: STRAIN: Hep-III
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       YFDLRLKGDK 13
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STREET: lov
-mv: Chicago
                                                 FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          60601
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCT-US95-02478-4
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APPLICANT: HINCH, TORU
APPLICANT: HINCH, TOSHIKI
APPLICANT: MINE, TOSHIKI
APPLICANT: MINE, TOSHIKI
APPLICANT: TYSON, ROBERT HUW
APPLICANT: TYSON, ROBERT HUW
APPLICANT: PAGE, ANTHONY MILES JOHN
TITLE OF INVENTION: 1-PHOSPHOTRANSFERASE ORIGINATING FROM PLANT, RECOMBINANT VECTOF
TITLE OF INVENTION: 3-PHOSPHOTRANSFERASE ORIGINATING FROM PLANT, RECOMBINANT VECTOF
TITLE OF INVENTION: 1-PHOSPHOTRANSFERASE ORIGINATING SUGAR CONTENT IN PLANT CELLS UNDEF
TUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: BIRCH, STEWART, KOLASCH AND BIRCH, LLP
STREET: PO BOX 747
CITY: FALL CHURCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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                                                        Gaps
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                                                      1; Indels
Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: FALL CHURCH
STATE: VA
COUNTRY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/416,870C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 32, Application US/08232539D
; Patent No. 5965709
; GENERAL INFORMATION:
; APPLICANT: Presta, Leonard G.
; APPLICANT: Jardieu, Paula M.
; TITLE OF INVENTION: IGE Antagonists
; NUMBER OF SEQUENCES: 60
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) OTHER INFORMATION: Mutant sequence substituted in place of IgE Fc(epsilon)3 fragment US-08-466-163B-50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.0%; Score 32; DB 4; Length 9; 100.0%; Pred. No. 3e+05; Cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 50, Application US/08466163B

Patent No. 6329509

GENERAL INFORMATION:

APPLICANT: Jardieu, Paula M.

APPLICANT: Jardieu, Paula M.

APPLICANT: Presta, Leonard G.

TITLE OF INVENTION: Immunoglobulin Variants

FILE REFERENCE: P0718P2C1D1

CURRENT APPLICATION NUMBER: US/08/466,163B

CURRENT FILING DATE: 1995-06-06

PRIOR PPLICATION NUMBER: US 08/405,617

PRIOR APPLICATION NUMBER: US 08/185,899

PRIOR FILING DATE: 1995-03-15

PRIOR FILING DATE: 1992-05-07

PRIOR FILING DATE: 1992-05-07

PRIOR FILING DATE: 1992-05-07

PRIOR FILING DATE: 1991-08-14

NUMBER OF SEQ ID NOS: 64

SEQ ID NOS: 64
               FILING DATE: 15-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/185899
FILING DATE: 26-JAN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/879495
FILING DATE: 07-MAY-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/744768
FILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: SVODOGA, Craig G.
REGISTRATION NUMBER: 39,044
REFERENCE/DOCKET NUMBER: P0718P;
TELEPHONE: 650/225-1489
TELEFAX: 650/952-9881
TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 50:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5; Conservative
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Best Local Similarity
Matches 5; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
TOPOLOGY: Linear
US-08-466-151-50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         153 WLDGK 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       153 WLDGK 157
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US-08-466-163B-50
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US-09-802-096-50
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                                                                                                                                                                 CUUNITRI: CCC
ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/232,539D
FILING DATE: 21-Apr-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/178583
FILING DATE: 07-JAN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/744768
FILING DATE: 14-AuG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,044
REFERENCE/DOCKET UNMBER: 90718P3
TELEPHONE: 650/952-9881
INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
TENGTH OF TELEPACE OF TENGTH OF TENGTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Immunoglobulin Variants
NUMBER OF SEQUENCES: 65
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
COUNTRY: USA
ZIP: 94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-466-151-50
; Sequence 50, Application US/08466151
; Patent No. 6037453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/466163
FILING DATE: 06-Jun-1995
ADDRESSEE: GELL.
STREET: 1 DNA Way
CITY: South San Francisco
                               Genentech, Inc.
  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: Amino Acid
TOPOLOGY: Linear
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                                                   3.0%; Score 32; DB 2; Length 10; 100.0%; Pred. No. 2.4e+03; iive 0; Mismatches 0; Indels
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                                                                                                                                                                             RESULT 41
US-08-466-151-17
; Sequence 17, Application US/08466151
; Patent No. 6037453
; GENERAL INFORMATION:
; APPLICANT: Jardieu, Paula M.
; APPLICANT: Presta, Leonard G.
; TITLE OF INVENTION: Immunoglobulin Variants
NUMBER OF SEQUENCES: 65
; CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,151
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CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION NUMBER: 08/466163
FILING DATE: 06-Jun-1995
APPLICATION NUMBER: 08/405617
FILING DATE: 15-MAR-1995
PILING DATE: 15-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/185899
FILING DATE: 26-JAN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/879495
FILING DATE: 07-MAY-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/879495
FILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,044
REFERENCE/DOCKET NUMBER: P0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1489
                                                   Query Match
Best Local Similarity 100.
Matches 5; Conservative
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US-08-466-151-17
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US-08-464-025A-9
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                                                                      APPLICANT: Dardieu, Paula M.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Method of Preventing the Onset of Allergic Disorders (as amended TITLE OF INVENTION: Method of Preventing the Onset of Allergic Disorders (as amended FILE REFERENCE: P0718P2C3US
FILE REFERENCE: P0718P2C3US
CURRENT FILING DATE: 2001-03-08
FRIOR FILING DATE: 1995-03-15
FRIOR FILING DATE: 1995-03-15
FRIOR FILING DATE: 1994-01-26
FRIOR FILING DATE: 1992-08-14
FRIOR FILING DATE: 1992-08-14
FRIOR FILING DATE: 1992-08-14
FRIOR FILING DATE: 1991-08-14
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US-09-802-096-50
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100.0%; Pred. No. 3e+05;
ative 0; Mismatches 0; Indels
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ZIP: 94080

COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/464,025A
FILING DATE: 05-Jun-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-464-025A-9

Sequence 9, Application US/08464025A

Patent No. 5994514

GENERAL INFORMATION:

APPLICANT: Jardieu et al.

TITLE OF INVENTION: IMMUNOGLOBULIN VARIANTS

NUMBER OF SEQUENCES: 27

CORRESPONDENCE ADDRESS:

ADDRESSEE: Genentech, Inc.

STREET: 1 DNA Way

CITY: South San Francisco

STATE: California

COUNTRY: USA
Sequence 50, Application US/09802096
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CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,044
REFERENCE/DOCKET NUMBER: P0716
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Artificial sequence
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INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS;
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Best Local Similarity 100.
Matches 5, Conservative
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                        Patent No. 6685939
GENERAL INFORMATION:
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TOPOLOGY:
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Gaps

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; OTHER INFORMATION: Mutant sequence substituted in place of IgE Fc(epsilon)3 fragment
US-08-466-163B-46
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APPLICANT: ALIBHAI, MURTAZA F.
APPLICANT: ASTWOOD, JAMES D.
APPLICANT: SAMPSON, HUGH A.
APPLICANT: MCWHERTER, CHARLES A.
TITLE OF INVENTION: PREPARATION OF DEALLERGENIZED PROTEINS AND PERMUTEINS FILE REFERENCE: 11899.0217.NPUSOO (MOBT217)
CURRENT APPLICATION NUMBER: US/09/755,630B
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Best Local Similarity 100.0%; Pred. No. 2.4e+03;
Matches 5; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 46, Application US/08466163B; Batent No. 6329509; GENERAL INFORMATION:
APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Immunoglobulin Variants; FILE REFERENCE: P0718P2C1D1
CURRENT APPLICATION NUMBER: US/08/466,163B; CURRENT FILING DATE: 1995-06-06; PRIOR APPLICATION NUMBER: US 08/185,899; PRIOR FILING DATE: 1995-03-15; PRIOR APPLICATION NUMBER: US 07/879,495; PRIOR FILING DATE: 1992-05-07; PRIOR FILING DATE: 1992-05-07; PRIOR FILING DATE: 1991-08-14; NUMBER OF SEQ ID NOS: 64; SEQ ID NO 46; LENGTH: 10
PRIOR FILLING DATE: 1995-03-15
PRIOR APPLICATION NUMBER: US 08/185,899
PRIOR FILLING DATE: 1994-01-26
PRIOR PILLING DATE: 1994-01-26
PRIOR PILLING DATE: 1992-05-07
PRIOR PILLING DATE: 1992-05-07
PRIOR PILLING DATE: 1991-08-14
NUMBER OF SEQ ID NOS: 64
SEQ ID NO 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 33, Application US/09755630B Patent No. 6639054
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                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.
Matches 5; Conservative
                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
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US-08-466-163B-46
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US-09-755-630B-33
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                                                                                                                                                                                                                                                                                                                                       ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,151
                                                                      GENERAL INFORMATION:
APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Immunoglobulin Variants
NUMBER OF SEQUENCES: 65
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17, Application US/08466163B; Patent No. 6329509; GENERAL INFORMATION:
APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Immunoglobulin Variants; FILE REFERENCE: P0718P2CID1
CURRENT APPLICATION NUMBER: US/08/466,163B; CURRENT FILING DATE: 1995-06-06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 08/466163
FILING DATE: 06-Jun-1995
APPLICATION NUMBER: 08/405617
FILING DATE: 15-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/185899
FILING DATE: 26-JAN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/879495
FILING DATE: 07-MAY-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/744768
FILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,044
REFERENCE/DOCKET NUMBER: 90718P;
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/952-9881
INFORMATION FOR SEQ ID NO: 46:
                           Sequence 46, Application US/08466151 Patent No. 6037453
                                                                                                                                                                                                                                             STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.0
Matches 5, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: Linear
US-08-466-151-46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: Amino Acid
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US-08-466-163B-17
         US-08-466-151-46
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Gaps

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CURRENT FILING DATE: 2001-01-05

PRIOR APPLICATION NUMBER: US 60/174,669

PRIOR FILING DATE: 2000-01-06

NUMBER OF SEQ ID NOS: 295

SOFTWARE: Patentin version 3.0

SOFTWARE: PRT

TYPE: PRT

CORGANISM: Artificial Sequence

FEATURE:

CORGANISM: Artificial Sequence

FEATURE:

CORGANISM: Artificial Sequence

FEATURE:

MS-09-755-630B-33

Query Match

Best Local Similarity 75.0%; Pred. No. 2.4e+03;

Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps

QY

201 KPFAIAKE 208

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Search completed: August 30, 2004, 10:57:21 Job time : 83.2162 secs

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                            Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                         Aar48203
Aaw67692
Aab82292
Abu57332
Add88405
Aaq918183
Aay43067
Aaw82306
Aaw82306
                                                                                                                       Aar92146 |
Abb74381 |
Aar35255 |
Aar56723 |
Aar91471 |
Aaw25253 |
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Pred. No. 1.7e+06;
0; Mismatches 1; Indels
                                                                                                                                                                                              ALIGNMENTS
                                                                                                                                                                                                                                                                                            Human cyclophilin B peptide fragment #42
                                                                                                                       AAR92146
ABB74381
AAR35255
AAR56723
AAR91471
AAW25253
                                 AAW67692
AAB82292
ABU57332
ADD88405
AAD918183
AAP43067
AAW43067
AAW8306
AAW8306
AAW8306
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ilarity 87.5%;
Conservative C
WEI; 2000-116932/10.
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Best Local Similarity
Matches 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 9 AA;
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AAY69962;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Abg79075 Human Cyp
Abr84373 Human Cyp
Adc17701 Cyclophil
Adg89641 Class I H
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Add88394 Tumour vi
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Human cyc
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(without alignments)
38.898 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Aay69921 H
Aab46946 H
Aab46930 H
Aag68088 A
Abg79075 H
Abr84373 H
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Ado33999 B
Abg32837 E
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       GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                          Total number of hits satisfying chosen parameters:
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                                         - protein search, using sw model
                                                          October 21, 2004, 07:04:19
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AAY69960
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AAB46946
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AAG68088
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ADF66836
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AD033999
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ABG94771
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39
                                                                                                             1 KXHRVIKDX 9
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Match Length
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No.

Result

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This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                              Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                   Claim 4; Page 49; 64pp; Japanese.
                                                                                                (SUMU ) SUMITOMO PHARM CO LID (ITOH/) ITOH K.
                                                          99WO-JP003360
                                                                             98JP-00178449
                                                                                                                                                                                                                                                                                                    Best Local Similarity 87.5
Matches 7; Conservative
                                                                                                                                                  WPI; 2000-116932/10.
                                                                                                                                                                                                                                                                                                                                   1 KXHRVIKD
                                                                                                                                                                                                                                                                                                                                                    KFHRVIKD
                                                                                                                              Itoh K, Gomi S;
                                                                                                                                                                                                                                                                        Sequence 9 AA;
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EP1074267-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-JUL-1999;
                  WO9967288-A1
                                                          24-JUN-1999;
                                                                             25-JUN-1998;
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                                     29-DEC-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                           AAB46946;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL;
HLA antigen; diagnosis; tumour; therapy.
                                                                                                                               Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                         Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
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                                                                                                                                                                                                                        Met
                                                                                                                                                                                                                       Trp,
                                                                                                                                                                                                    'label= Phe, Tyr, Met, Trp
                                                                                                             Human cyclophilin B peptide fragment #40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human cyclophilin B peptide fragment #1.
                                                                                                                                                                                                                       /label= Phe, Leu, Ile,
                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 10; Page 59; 64pp; Japanese.
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                                                 AAY69960 standard; peptide; 9 AA.
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                                                                                                                                                                                                                                                                                                                          (SUMU ) SUMITOMO PHARM CO LTD
                                                                                                                                                                                                                                                                                   99WO-JP003360
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                                                                                         (first entry)
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Best Local Similarity
Matches 8; Conserv
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1 KYHRVIKD
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                                                                                         11-APR-2000
                                                                                                                                                               Homo sapiens
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ID AAY
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Gaps

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1; Indels Length 9;

92.3%; Score 36; DB 3; I 87.5%; Pred. No. 1.7e+06; ive 0; Mismatches 1;

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Use of antigenic proteins, peptides, interferon or their encoding DNA, in the manufacture of an agent for the induction of antigen-specific {\tt T}
                                                                                                                                                                                                                                                                            Tumor antigenic peptide; interferon; IFN; antigen-specific T cell. virucide; cytostatic; gene therapy; IFN-alpha; cytotoxic T cell; viral infectious disease; cyclophilin B; human.
                                                                                                                                                                                                               Human cyclophilin B derived tumor antigenic peptide SEQ ID 32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 19; 25pp; English
AAB46946 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yamaoka T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SUMU ) SUMITOMO PHARM CO LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-JUL-2000; 2000EP-00306263.
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                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-193144/20
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This invention describes the novel use of interferons (IFNs) or DNAs capable of expressing the interferons and/or antigenic proteins (AP), antigenic peptides derived from the proteins or DNAs capable of expressing the antigeners or peptides, in the manufacture of an agent for induction of antigen-specific T cells. The products of the invention have virucide and cytostatic activity and can be used for gene therapy or as inducers of antigen-specific T cells. The action of interferon-alpha (IFN-alpha) in a system for inducing specific cytotoxic T cell (CTL) by administering an antigenic peptide in an incomplete Freund's adjuvant (IFA) emulsion preparation form was evaluated. IFNs (or DNA encoding IFNs) are useful in the manufacture of a medicament for inducing antigen-specific T cells in an individual who has been administered with AP (or DNA encoding AP) or vice versa. The medicament is useful for the treatment or prophylaxis of a tumor or a viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                 infectious disease
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Sequence 9 AA;

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                    Gaps
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                    1; Indels
     92.3%; Score 36; DB 4; Length 9; 87.5%; Pred. No. 1.7e+06;
                    0; Mismatches
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1 KXHRVIKD 8

ထ 1 KYHRVIKD

AAB46930

AAB46930 standard; peptide; 9 AA.

AAB46930;

(first entry) 04-MAY-2001 Human cyclophilin B derived tumor antigenic peptide SEQ ID 16.

Tumor antigenic peptide, interferon, IFN; antigen-specific T cell virucide, cytostatic, gene therapy; IFN-alpha, cytotoxic T cell; viral infectious disease; cyclophilin B; human.

Homo sapiens

EP1074267-A1

24-JUL-2000; 2000EP-00306263.

99JP-00207687 22-JUL-1999; (SUMU) SUMITOMO PHARM CO LTD

Gotoh M, rakasu H,

WPI; 2001-193144/20

Use of antigenic proteins, peptides, interferon or their encoding DNA, in the manufacture of an agent for the induction of antigen-specific ${\tt T}$

Disclosure; Page 15; 25pp; English.

This invention describes the novel use of interferons (IFNs) or DNAs capable of expressing the interferons and/or antigenic proteins (AP), antigenic peptides derived from the proteins or DNAs capable of expressing the antigenic proteins or peptides, in the manufacture of an agent for induction of antigen-specific T cells. The products of the invention have virucide and cytostatic activity and can be used for gene therapy or as inducers of antigen-specific T cells. The action of interferon-alpha (IFN-alpha) in a system for inducing specific cytotoxic T cell (CTL) by administering an antigenic peptide in an incomplete

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Gaps

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92.3%; Score 36; DB 4; Length 9; 87.5%; Pred. No. 1.7e+06; tive 0; Mismatches 1; Indels

Query Match Best Local Similarity 87.5% Lagaryative

Sequence 9 AA;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tumour specific cytotoxic I lymphocytes (CTL). The peptides recognising cancer cells with tumour specific CTLs are selected from: (1) peptides of sequences (AAG68066 to AAG68069); (2) peptides containing the above mentioned sequences; (3) peptides having 70 % or more of homogeneity with the above mentioned sequences; and (4) peptides with one or more deleted, substituted, added or inserted amino acid(s) of the above mentioned sequences, particularly those having recognising property due to HLA-A2402 binding CTL, especially having at least 5 amino acids, used for medicine, particularly anticancer agents, derived from antitumour antigenic peptides of 1ck, src family, SART-1, SART-3 or cyclophilin B genes. The antitumour peptides have cytostatic activities. The peptides are used for the treatment of cancer. The peptides cause activation of the present invariant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New peptides for recognizing cancer cells with tumor specific cytotoxic T lymphocytes and for treating cancer.
         (or
Freund's adjuvant (IFA) emulsion preparation form was evaluated. IFNs (c DNA encoding IFNs) are useful in the manufacture of a medicament for inducing antigen-specific T cells in an individual who has been administered with AP (or DNA encoding AP) or vice versa. The medicament is useful for the treatment or prophylaxis of a tumor or a viral infectious disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antitumour; cancer; cancer cell recognition; antigenic; CTL; lck; src; tumour specific cytotoxic T lymphocyte; anticancer; SART-1; SART-3; cyclophilin B gene; HLA-A2402.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes peptides recognising cancer cells with
                                                                                                                                                                                                               Gaps
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                                                                                                                                                                        92.3%; Score 36; DB 4; Length 9; 87.5%; Pred. No. 1.7e+06; iive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antitumour peptide cyclophilin B 84-92.
                                                                                                                                                                                                                                                                                                                                                                                 AAG68088 standard; peptide; 9 AA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-DEC-2000; 2000JP-00393047.
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                                                                                                                                                                                           Local Similarity 87.5
nes 7; Conservative
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                                                                                                                                                                                                                                                   1 KXHRVIKD 8
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                                                                                                                                          Sequence 9 AA;
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cell and a cell penetrating peptide (CPP) associated with an antigen or antibody. Also included are (1) a vaccine comprising (I), CPP associated with an antigen, and a pharmaceutically acceptable carrier and (2) preparing a composition for a disease, by providing (I) and CPP associated with an antigen for disease, and introducing the antigen-sessociated CPP to (I), where antigen enters into the cell. The antigens or infiltrating lymphocytes (TIL) of HiA (human leukocyte antigen) class I or II. The composition is useful for enhancing immunity in an animal to a disease, by administering a mature dendritic cell comprising CPP associated with an antigen to disease, to the animal, such that following the administration, animal is protected from disease, where the animal comprises both CD4+ and CD8+ T cells. It is also useful for treating a disease (c) cancer, tumour, melanoma, lymphoma, uterine cancer, non-Eddgkin's lymphoma, leukaemia, Hodgkin's lymphoma, uterine cancer, cervical cancer, bladder cancer, kidney cancer, adenocarcinoma, cradiation, chemotherapy or gene therapy. The administration of (I), rediation, chemotherapy or gene therapy. The administration of (I), rediation, chemotherapy or gene therapy. The administration of (I), the cancer treatment. The present sequence is a tumour antigen derived epitope for inclusion in the composition of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a composition (I) comprising an immune effector
                                                                                                                                                                                                                                                                                                             Cell penetrating peptide; cancer; tumour; melanoma; thymoma; antigen;
                                                                                                                                                                                                                                                                                                                              lymphoma; sarcoma; lung cancer; non-Hodgkin's lymphoma; leukaemia; Hodgkin's lymphoma; uterine cancer; cervical cancer; bladder cancer; kidney cancer; adenocarcinoma; breast cancer; prostate cancer; ovarian cancer; pancreatic cancer; epitope; vaccine; dendritic cell; tumour infiltrating lymphocyte; TIL; human leukocyte antigen; HLA; cytostatic; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel composition for treating a disease in an animal, comprises an immune effector cell and cell penetrating peptide associated with an antigen or antibody.
                                                                                                                                                                                                                                                                   Human Cyp-B class I HLA widely expressed antigen peptide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 17; 61pp; English
                                                                                                                                             ABG79075 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BAYU ) BAYLOR COLLEGE MEDICINE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-FEB-2002; 2002WO-US005212.
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                                                                                                                                                                                                                            (first entry)
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KXHRVIKD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                            15-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-AUG-2002
                                                                                                                                                                                     ABG79075;
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                                                                                                     RESULT 7
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The invention relates to a method for the detection of antigen specific T-cells in a blood sample involving the use of a plurality of antigenic peptides. The method comprises sampling of peripheral blood monocytes; stimulation of the collected peripheral blood monocytes with antigens without direct use of antigen presenting cells; and detection of T-cells specific to the antigen in the stimulated monocytes. The method is particularly used for the detection of cancer as it can be used in semi-quantitative determination of cancer specific T-cells. It can also be used for cancer vaccine therapy for patients with cervical or prostate cancer. The method can additionally be used to monitor of cellular immunity and cancer immune therapy by detection of specific T-cell frequencies. Sequences ABR84367-ABR84380 represent HLA-A24 (human
                                                                                                                                                                                                                                            Antigen specific T-cell; detection; diagnosis; cancer specific T-cell; cancer; tumour; cervical cancer; prostate cancer; cellular immunity; immune therapy; cytostatic; immunostimulant; vaccine; antigenic peptide; human; human leukocyte antigen; HLA-A24 epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A detection method of antigen specific T-cells, comprises the use of plural antigenic peptides, useful in semi-quantitative determination of cancer specific T-cell frequencies and for monitoring cellular immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                eukocyte antigen) peptides of human origin used in an example from the
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                                                                                                                                                                                                          Human CypB HLA-A24 epitope, SEQ ID NO:23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 8; Page 10; 18pp; Japanese.
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                                                                                             ABR84373 standard; peptide; 9 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-NOV-2000; 2000JP-00345094
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87.5%;
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Best Local Similarity
KFHRVIKD
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                                                                                                                                                                                                                                                                                                                                            Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                  18-DEC-2002
                                                                                                                                 ABR84373;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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ADC17701
                                                                          ABR84373
                                                         RESULT
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Length 9;

Score 36; DB 5;] Pred. No. 1.7e+06;

92.3%;

Query Match Best Local Similarity

us-09-720-469a-37.rag

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Identifying a cell that differentiates into a metastatic cancer cell, useful for preventing metastatic cancer, comprises identifying a mutated fibronectin in the cell.
                                                                                                                                                                                                                                                                                        The invention comprises a method for identifying a cell that will differentiate into a metastatic cancer cell, the method involves identifying a mutated fibronectin in the cell. The method of the invention is useful for preventing metastatic cancer. The present amino acid sequence represents a Class I HLA-restricted widely expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 36; DB 8; Length 9;
Pred. No. 1.7e+06;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                               Disclosure; SEQ ID NO 84; 137pp; English.
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                                                                                            (BAYU ) BAYLOR COLLEGE MEDICINE
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27-MAR-2001; 2001US-0278761P.
09-JUL-2001; 2001US-0303396P.
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                             28-MAY-2003; 2003WO-US016736.
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                                                               2002US-0383530P
                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 92.3%;
Best Local Similarity 87.5%;
Matches 7; Conservative
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                                                                                                                                                               WPI; 2004-035134/03
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                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polyoma virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-SEP-1998;
                                                               28-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-AUG-2002.
04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                               antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to allergic reaction suppressors, comprising a peptide which originates from the same antigenic substance as the antigenic substance inducing the (peptide-originated) allergic reaction, and containing an epitope different from the epitope participating in the induction of the allergic reaction and yet dos not induce the allergic reaction. The desensitizers or allergic reaction suppressors are useful for treating and preventing type I allergic diseases and in cancer vaccines for preventing or treating cancer. The allergic reaction suppressors are also useful for suppressing an allergic reaction or reducing immunoglobulin E antibody production, and for desensitization to antigens. This sequence represents a peptide used in the invention and corresponds to amino acids 84-92 of the cyclophilin B protein.
                                                                               vaccine; allergic reaction suppressor; antigen;
immunoglobulin E antibody; desensitization;
                                                                                                                                                                                                                                                                                                                                                                                                                Desensitizers or allergic reaction suppressors containing peptides originating from the same antigenic substances as reaction-inducing substances, useful for preventing or treating type I allergic diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 metastatic cancer cell differentiation; mutated fibronectin; metastatic cancer; class I HLA-restricted; widely antigen.
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                                                Cyclophilin B protein amino acids 84-92.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 8; SEQ ID NO 1; 49pp; Japanese.
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                                                                                                                                                                                                                                                28-AUG-2002; 2002WO-JP008641.
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                                                                            cytostatic; antiallergic;
allergy; epitope; cancer;
cyclophilin B.
                 (first entry)
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Best Local Similarity 8/...
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                 18-DEC-2003
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                                                                                                                                                Synthetic.
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Replikin; glioma; malignin; anthrax; small pox; antiviral; antibacterial; 3-point-recognition; vaccine; immunogen; Lethal Factor Protein pX01-107; Surface Antigen S Precursor Protein.
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Replikin sequence identiied by the 3-point recognition method #44.
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                                                     The Inversion relates to located six to ten residues from a second lysine residue, at least one histidine residue, and at least 6% including at least one lysine residue located six to ten residues from a second lysine residue, at least one histidine residue, and at least 6% including at least one histidine residue, and at least 6% least one the repplikin sequence isolated from glioblastoma cells malignin peptide which is of non-human origin. Also included are anti-replikin antibodies (or antibody cocktail), a method of stimulating the immune system of a subject to produce antibodies to amount of at least one or more Bacillus anthracis or small pox comprising administering an effective mount of at least one or more Bacillus anthracis or small pox replikin antisense nucleic acid molecule complementary to a mRNA encoding a Bacillus anthracis polypeptide (e.g. Anthrax Lethal Factor Protein pX01-107) comprising a replikin sequence, an antisense nucleic acid molecule complementary to a mRNA encoding a Small Pox Virus polypeptide (e.g. Surface Antigen S Precursor Protein comprising a replikin sequence. The peptides are useful for simulating the immune system of a subject to produce antibodies that bind specifically to Anthrax or small pox prevention of produce antibodies that can be used in the treatment and/or prevention of anthrax or small pox. The peptides are useful for generating antibodies that can be used in the treatment and or prevention of anthrax or small pox. The peptides are useful for generating anthrax or small pox. The peptides are useful for structural similarity with the replikin sequence isolated from structural similarity with the replikin sequence isolated from a school to a produce of structural similarity with the replikin sequence isolated from system and sharing to the structural similarity with the replikin sequence isolated from a school to the structural similarity with the replikin sequence isolated by the structure isolated from the structure isolated by the structure isolated by the stru
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                                               pox virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Replikin sequence identiied by the 3-point recognition method #45.
                                               and small
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 28; DB 5; Length 9;
Pred. No. 1.7e+06;
1; Mismatches 2; Indels
                                              invention relates to isolated Bacillus anthracis
              Disclosure; Page 3; 16pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABG94772 standard; peptide; 9 AA.
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27-MAR-2001; 2001US-0278761P.
09-JUL-2001; 2001US-0303396P.
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62.5%;
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Best Local Similarity 62.5
Matches 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 KXHRVIKD 8
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BOGOCH E S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polyoma virus.
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The invention relates to isolated Bacillus anthracis and small pox virus comptides, also known as replikins, comprising 7 to 50 amino acids including at least one lysine residue located six to ten residues from a second lysine residue, at least one histidine residue, and at least 6% including at least one histidine residue, and at least 6% second lysine residues. The peptides were identified using the 3-point-recognition method and based on the replikin sequence isolated from glioblastoma cells malignin peptide which is of non-human origin. Also included are anti-replikin antibodies (or antibody cocktail), a method of stimulating the immune system of a subject to produce antibodies to said molecule complementary to a manaly pox comprising administering an effective amount of at least one or more Bacillus anthracis or small pox replikin anthracis polypeptide (e.g. Anthrax Lethal Factor Protein pX01-07) complementary to a manaly encoding a Small Pox virus polypeptide (e.g. Anthrax Lethal Factor Protein pX01-07) complementary to a manaly encoding a Small Pox virus polypeptide (e.g. Surface Antigen S Precursor Protein comprising a replikin sequence. The peptides are useful for stimulating the immune system of a subject to produce antibodies that bind specifically to Anthrax or small pox produce antibodies that bind specifically to Anthrax or small pox of anthrax or small pox. The peptides are useful for sequence is a replikin sequence isolated by the 3-point-recognition system and sharing structural similarity with the replikin sequence isolated from sequence isolated from the replikin sequence isolated from the sequence isolated by the 3-point solated from the sequence isolated by the 3-point solated from the sequence isolated by the 3-point solated from the sequence isolated by the 3-point sequence is
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Isolated Bacilius anthracis and small pox virus peptides (I), also known as replikins, useful for formulating vaccines for treating anthrax or small pox, comprise 7 to 50 amino acids.
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                                                                                                                                                                             Disclosure; Page 3; 16pp; English.
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62.5%;
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09-JUL-2001; 2001US-0303396P.
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Best Local Similarity
Matches 5; Conserv
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KQHRELKD
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                                                                                                         The invention describes an isolated replikin or recognin peptide (I) comprising 7-50 amino acids including at least one lysine residue located 6 to 10 residues from a second lysine residue, at least one histidine residue and at least 6% lysine residues. The peptides are useful for stimulating the immune system, and for targeting, labelling or destroying replikin-containing organisms. The computer-readable mediums are useful for identifying a protein or peptide containing a replikin or recognin sequence. This is the amino acid sequence of a replikin peptide identified in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New replikins or recognin peptides, useful for stimulating the immune system, and for targeting, labeling or destroying replikin-containing
            likins or recognin peptides, useful for stimulating the immune and for targeting, labeling or destroying replikin-containing
                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                  .;
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                                                                                                                                                                                                                                                                                                               71.8%; Score 28; DB 6; Length 9; 62.5%; Pred. No. 1.7e+06; iive 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Replikin; immunostimulant; vaccine; recognin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABU57322 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 79; Page 3; 17pp; English.
                                                                              Claim 78; Page 3; 17pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Replikin peptide sequence #62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-OCT-2001; 2001US-00984057.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-MAR-2001; 2001US-0278761P.
09-JUL-2001; 2001US-0303396P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                Bogoch ES;
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                                                                                                                                                                                                                                                                                                                                                                                  1 KXHRVIKD 8
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(BOGO/) BOGOCH E S.
                                                                                                                                                                                                                                                                                                                                                                                                                   KOHRELKD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2002151677-A1
                                                                                                                                                                                                                                                                                  Sequence 9 AA;
              replikins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-OCT-2002
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                                               organisms.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an isolated Plasmodium falciparum peptide or an influenza virus peptide. The composition and methods are useful in diagnosing, preventing or treating influenza virus infection or malaria. Also disclosed is an influenza virus vaccine comprising at least one isolated Replikin present in the haemagglutinin protein of an emerging strain of influenza virus and a pharmaceutical carrier and/or adjuvant and a malaria vaccine comprising at least one isolated P. falciparum Replikin and a pharmaceutical carrier. The present sequence represents the amino acid sequence of a Replikin sequence.
                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                      vaccine; gene therapy; malaria; influenza; Replikin; haemagglutinin.
                                                              ..
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                        Score 28; DB 6; Length 9; Pred. No. 1.7e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          71.8%; Score 28; DB 7; Length 9; 62.5%; Pred. No. 1.7e+06; tive 1; Mismatches 2; Indels
                                                            1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; SEQ ID NO 61; 136pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  completed: October 21, 2004, 07:20:58
                                                                                                                                                                                 ADD88394 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                       27-MAR-2001; 2001US-0278761P.
09-JUL-2001; 2001US-0303396P.
26-OCT-2001; 2001US-00984057.
                                                                                                                                                                                                                                                                                                                                                                                                26-MAR-2002; 2002US-00105232
                                    71.8%;
62.5%;
                                                                                                                                                                                                                                                              Tumour viral Replikin #14.
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                                                              Conservative
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Best Local Similarity
                                    Query Match
Best Local Similarity
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                                                                                        1 KXHRVIKD
                                                                                                                 1 KOHRELKD
                                                                                                                                                                                                                                                                                                                                            US2003180328-A1.
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          Sequence 9 AA;
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                                                                                                                                                                                                                                                                                                                  Polyomavirus
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                                                                                                                                                                                                           ADD88394;
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; Sequence 1, Application US/08471286
; Patent No. 5866688
; GENERAL INFORMATION:
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TYPE: amino acid
STRANDEDNESS: single
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Best Local Similarity 83.3
Matches 5; Conservative
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CLASSIFICATION: 530
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US-08-471-286-1
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                                                            October 21, 2004, 07:13:10 ; Search time 21.5 Seconds (without alignments) 27.761 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                Issued_Patents_AA:*
1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-471-286-1

US-08-378-761A-68

US-08-485-286-68

US-10-037-311B-16

US-09-418-780A-22

US-09-418-780A-22

US-09-418-780A-22

US-07-963-321-29

US-08-290-641-29

US-08-290-641-29

US-08-390-641-29

US-08-390-641-29

US-08-391-391

US-09-344-040C-114

US-09-344-040C-114

US-09-333-28

US-09-331-28

US-09-331-28

US-09-331-28

US-09-331-57

US-07-778-233B-28

US-07-963-321-28

US-07-963-321-57

US-08-290-641-28

US-08-290-641-57

US-08-548-540-28
                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCT-US96-09809-28
PCT-US96-09809-57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-057-363C-26
                                                                                                                                                                                 478139 segs, 66318000 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                 SUMMARIES
                                                                                                                                                                                                                                                        Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                         - protein search, using sw model
                                                                                                                                                 BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                     US-09-720-469A-37
39
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query
Match Length DB
                                                                                                                            1 KXHRVIKDX 9
                                                                                                                                                                                                                         Minimum DB seq length: 0 Maximum DB seq length: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Title:
Perfect score:
                                                                                                                                                  Scoring table:
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                                                                                                                              Sequence:
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                                                              Run on:
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  Sequence 26, Appl
Sequence 46, Appl
Sequence 46, Appl
Sequence 48, Appl
Sequence 60, Appl
Sequence 314, App
Sequence 314, App
Sequence 314, App
Sequence 205, Appl
Sequence 605, Appl
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0
                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Kim, Hyesook
APPLICANT: Charnecki, Jonathan
APPLICANT: Putt, David A.
APPLICANT: Kim, Edward Y.
TITLE OF INVENTION: PRODUCTION OF ANTI-PEPTIDE ANTIBODIES
TITLE OF INVENTION: AGAINST CYTOCHROME P450
NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Michigan
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 48099-4390
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/471,286
US-09-265-107-26
US-08-893-534A-46
US-08-996-679-46
US-09-115-395-76
US-09-507-102-46
US-09-250-059-48
US-09-250-059-48
US-09-357-717-30
US-09-357-717-30
US-09-248-074-48
US-09-244-782-48
US-09-244-782-48
US-09-234-395-314
US-09-305-928-314
US-09-305-928-314
US-08-56-597-122
US-08-856-597-122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  69.2%; Score 27; DB 2;
83.3%; Pred. No. 17;
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:

NAME: Kohn, Kenneth I.

REGISTRATION NUMBER: 30,955

REFERENCE/DOCKET NUMBER: P-301 (Oxford)

TELECOMMUNICATION INFORMATION:

TELEPHONE: (810) 689-3500

TELEFAX: (810) 689-4071

INFORMATION FOR SEQ ID NO: 1:

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                      US-08-742-243-6
                                                                                                                                                                                                                                                            ALIGNMENTS
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64.1%; Score 25; DB 4; Length 12; 66.7%; Pred. No. 45; 0; Indels :ive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 25; DB 1; Length 11;
Pred. No. 41;
1; Mismatches 2; Indels
ZIP: 46268
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn PC-DOS/MS-DOS
SOFTWARE: PatentIn PATA:
APPLICATION NUMBER: US/08/485,286
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/378761
FILING DATE:
CLASSIFICATION NUMBER: 33651
FILING DATE: 26-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: BORUCKI, ANDREA T
REGISTRATION NUMBER: 33651
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF 68:
TELECOMMUNICATION OF 68:
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Sequence 16, Application US/10037311B

Patent No. 6747190

GENERAL INFORMATION:

TITLE OF INVENTION: XYLOGLUCAN FUCOSYLTRANSFERASES

FILE REFERENCE: MS00-001C2

CURRENT APPLICATION NUMBER: US/10/037,311B

CURRENT FILING DATE: 2001-11-09

PRIOR PILING DATE: 1999-01-28

NUMBER OF SEQ ID NOS: 17

SOFTWARE: Patentin version 3.1

SOFTWARE: Patentin version 3.1
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Patent No. 618076B
GENERAL INFORMATION:
APPLICANT: MAERTENS, GEERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                64.1%;
62.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STUYVER, LIEVEN
                                                                                                                                                                                                                                                                                                                                                                                                                                        11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 , MOLECULE TYPE: protein US-08-485-286-68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: Cross-species
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: si:
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Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
Matches 5; Conserv
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HOVIRD 11
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                                                                            Sequence 68, Application US/08378761A
Sequence 68, Application US/08378761A
Patent No. 5635384
GENERAL INFORMATION:
APPLICANT: WALSH, TERENCE A
APPLICANT: WORGAN, ALICE ER
TITLE OF INVENTION: REBOSOME-INACTIVATING PROTEINS, INACTIVE
TITLE OF INVENTION: USING
NUMBER OF SEQUENCES: 81
CORRESPONDENCE ADDRESS:
ADDRESSEE: ANDREA T. BORUCKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: WALSH, TERENCE A
APPLICANT: WALSH, TERENCE A
APPLICANT: WORGAN, ALICE ER
TITLE OF INVENTION: RIBOSOME-INACTIVATING PROTEINS, INACTIVE
TITLE OF INVENTION: USING
NUMBER OF SEQUENCES: 81
CORRESPONDENCE ADDRESS:
ADDRESSEE: ANDREA T: BORUCKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 46268
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: DC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/378,761A
FILING DATE: 26-JAN-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                    STREET: 9330 ZIONSVILLE ROAD CITY: INDIANAPOLIS STATE: IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY AGENT INFORMATION:
NAME: BORUCKI, ANDREA T
REGISTRATION NUMBER: 33651
REFERENCE/DOCKET NUMBER: 3827
TELECOMMUNICATION INFORMATION:
TELEPHONE: (317) 337-4846
INFORMATION FOR SEQ ID NO: 68:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               : 9330 ZIONSVILLE ROAD INDIANAPOLIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   64.18;
62.58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 62.5
Teches 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 KXHRVIKD 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: si:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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                                                                    US-08-378-761A-68
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US-08-485-286-68
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Patent No. 5270170
GENERAL INFORMATION:
APPLICANT: Schatz, Peter J.
APPLICANT: Cull, Millard G.
APPLICANT: Miller, Jeff F.
TITLE OF INVENTION: Peptide Library and Screening Method NUMBER OF SEQUENCES: 78
CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Db 2,
; 1.8e+02;
... 0; Indels
                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     56.4%; Score 22; DB 4; Length 11; 60.0%; Pred. No. 1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/778,233B
FILING DATE: 19911016
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REPERENCE/DOCKET NUMBER: 11509-50
TELECOMMUNICATION INFORMATION:
TELEFAN: 415-326-2400
TELEFAN: 415-326-2400
TELEFAN: 415-326-2422
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
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0
                                                                                                                                                                                                                                                        GENERAL INCORMATION:
GENERAL INFORMATION:
APPLICANT: JONES, Michael H.
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATOR
FILE REFERENCE: 06501-043001
CURRENT APPLICATION NUMBER: US/09/418,780A
CURRENT FILING DATE: 1999-10-15
PRIOR APPLICATION NUMBER: PCT/JP98/01782
PRIOR FILING DATE: 1998-04-17
PRIOR FILING DATE: 1997-04-18
NUMBER OF SEQ ID NOS: 28
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 22
                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Mismatches
                                                                                                                                                                                            US-09-418-780A-22; Sequence 22, Application US/09418780A; Patent No. 6504009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TYPE: PRT
; ORGANISM: Drosophila melanogaster
US-09-418-780A-22
                5
             3; Conservative
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Best Local Similarity
Matches 3; Conserv
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2 HKIIK 6
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2 HKIIK 6
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US-07-778-233B-29
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              Matches
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           NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ·.
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Pred. No. 99;
2; Mismatches 2; Indels
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TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS
TITLE OF INVENTION: AND THEIR USE AS PROPHYLACTIC, THE
TITLE OF INVENTION: AGENTS
NUMBER OF SEQUENCES: 207
CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE & DURKEE
STREET: P.O. BOX 4433
CITY: HOUSTON
STATE: TEXAS
COUNTRY: USA
ZIP: 77210-4433
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 6.0 / ASCII text output
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/836,075A
FILING DATE: 21 Apr 1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/04155
FILING DATE: 23 Oct 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/04155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 14, Application US/09418780A; Patent No. 6504009; GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATOR; FILE REFERENCE: 06501-043001
CURRENT APPLICATION NUMBER: US/09/418,780A; CURRENT FILING DATE: 1999-10-15; PRIOR APPLICATION NUMBER: PCT/JP98/01782; PRIOR APPLICATION NUMBER: JAPAN 9/116402; PRIOR FILING DATE: 1997-04-18
NUMBER OF SEQ ID NOS: 28
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DAIA:
APPLICATION NUMBER: EP 94870166.9
FILING DATE: 21 Oct 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 95870076.7
FILING DATE: 28 Jun 1995
ATTORNEY/AGENT INFORMATION:
NAME: KAMMERER, PATRICIA A.
REGISTRATION NUMBER: 29,775
REFERENCE/DOCKET NUMBER: INNS:004
INFORMATION FOR SEQ ID NO: 204:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
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60.0%;
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Best Local Similarity 50.0
Matches 4, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT; ORGANISM: Homo sapiens
US-09-418-780A-14
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Best Local Similarity
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Patent No. 5338665

GENERAL INFORMATION:
APPLICANT: Schatz, Peter J.
APPLICANT: Cull, Millard G.
APPLICANT: Miller, Jeff F.
TITLE OF INVENTION: Peptide Library and Screening Method NUMBER OF SEQUENCES: 91
CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             56.4%; Score 22; DB 1; Length 12; 60.0%; Pred. No. 1.9e+02; ive 2; Mismatches 0; Indels
                                                                                                                                                                                          56.4%; Score 22; DB 1; Length 12; 60.0%; Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COFRAILING SISIEM: FC-DOS/MS-LOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/963,321
FILING DATE: 19921015
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/778,223
FILING DATE: 16-OCT-1991
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 11509-50-1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                   2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: One Market Plaza, Steuart CITY: San Francisco STATE: California COUNTRY: USA ZIP: 94105
COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 12 amino acids
TYPE: AMINO ACID
STRANDEDNESS: single
LENGTH: 12 amino acids
TYPE: AMINO ACID
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 60.0
Matches 3; Conservative
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Best Local Similarity 60.0
Matches 3; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide immediate source: CLONE: 26 4 0.8 US-07-963-321-29
                                                             TOPOLOGY: linear

MOLECULE TYPE: peptide

IMMEDIATE SOURCE:

CLONE: 26 4 0.8

US-07-778-233B-29
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HRVVR 10
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US-07-963-321-29
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/> Sequence 29, Application US/08290641
// Sequence 29, Application US/08290641
// Patent No. 5498530
// GENERAL INFORMATION:
// APPLICANT: Schatz, Peter J.
// APPLICANT: Miller, Jeff F.
// APPLICANT: Miller, Jeff F.
// TITLE OF INVENTION: Peptide Library and Screening Method NUMBER OF SEQUENCES: 91
// CORRESPONDENCE ADDRESS:
// ADDRESSEE: William M. Smith
// STREET: One Market Plaza, Steuart Tower, Suite 2000
// CITY: San Francisco
// STATE: California
// COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ASSOCIATION OF STATES APPLICATION US/08548540
; Sequence 29, Application US/08548540
; Patent No. 5733731
; GENERAL INFORMATION:
; APPLICANT: Cull, Millard G.
; APPLICANT: Miller, Willem P.C.
; APPLICANT: Stemmer, Willem P.C.
; APPLICANT: Gates, Christian M.
; TITLE OF INVENTION: Peptide Library and Screening Method
; NUMBER OF SEQUENCES: 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 22; DB 1; Length 12; Pred. No. 1.9e+02; 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 94105
ZIP: 94105
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/290,641
FILING DATE: 15-AUG-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/963,321
FILING DATE: 15-OCT-1992
APPLICATION NUMBER: US 07/778,223
FILING DATE: 16-OCT-1991
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
BERDDENGRY DOCUMENT AND THE STATION NUMBER: 30,223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11509-50-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 1150
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         56.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12 amino acids
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Best Local Similarity 60.00
There 3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: si
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US-08-290-641-29
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Sequence 93, Application US/08397633A;
Sequence 93, Application US/08397633A;
Patent No. 5773577;
GENERAL INFORMATION:
TITLE OF INVENTION: PRODUCTS COMPRISING SUBSTRATESCAPABLE
TITLE OF INVENTION: OF ENZYMATIC CROSS-LINKING
TITLE OF INVENTION: OF ENZYMATIC CROSS-LINKING
TITLE OF INVENTION: OF ENZYMATIC CROSS-LINKING
NUMBER OF SEQUENCES: 105
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: DISK PC COMPATIBLE
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Pred. No. 1.9e+02;
2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patent: E DOS/ES DOS
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                                              FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/548,540
FILING DATE: 26-OCT-1995
APPLICATION NUMBER: US 08/290,641
FILING DATE: 15-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/963,321
FILING DATE: 15-OCT-1992
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 16528J-001
TELECOMMUNICATION INFORMATION:
TELECHONE: 415-326-2400
TELECHONE: 415-326-2400
TELEFAX: 415-326-2400
APPLICATION NUMBER: PCT/US96/09809
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 93:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          56.48;
60.08;
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Best Local Similarity 60.0
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: peptide
IMMEDIATE SOURCE:
CLONE: 26 4 0.8
PCT-US96-09809-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6 HRVVR 10
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US-08-397-633A-93
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APPLICANT: Schatz, Peter J.
APPLICANT: Cull, Millard G.
APPLICANT: Miller, Jeff F.
APPLICANT: Stemmer, Willem P.C.
APPLICANT: Gates, Christian M.
TITLE OF INVENTION: Peptide Library and Screening Method NUMBER OF SEQUENCES: 162
CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          56.4%; Score 22; DB 1; Length 12; 60.0%; Pred. No. 1.9e+02; iive 2; Mismatches 0; Indels
                                          ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
CITY: San Francisco
                                                                                                                            CITY: SOLITION: CITY: CONTINUED CONTRY: CALIFORNIA COUNTRY: USA ZIP: 94105
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk COMPUTER: ISM PC COMPATIBLE COMPUTER: PATENTIN PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/548,540
FILING DATE: 26-OCT-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16528J-001240US
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APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/290,641
FILING DATE: 15-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/963,321
FILING DATE: 15-CCT-1992
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 16528J-0012
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRENCE/COST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 29: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 12 amino acids
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Best Local Similarity 60.0
Matches 3; Conservative
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   CORRESPONDENCE ADDRESS:
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STATE: California
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PCT-US96-09809-29
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Search completed: October 21, 2004, 07:29:28 Job time: 22.5 secs
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Sequence 72, Application US/09105839D

Patent No. 6287756

GENERAL INFORMATION:
APPLICANT: Tureci, Ozlem
APPLICANT: Chen, Yao-Tseng
APPLICANT: Sahin, Ugur
APPLICANT: Sahin, Ugur
APPLICANT: Old, Lloyd J
APPLICANT: Olf, Lloyd J
APPLICANT: Pfreundschuh, Michael
APPLICANT: Olf, Lloyd J
APPLICANT: Olf, Lloyd J
APPLICANT: Of INVENTION: Method for Determining Presence of Cancer In A Sample By Determin
TITLE OF INVENTION: Expression of an SSX gene
FILE REFERENCE: LUD 5556
CURRENT FILING DATE: 1998-06-26
CURRENT FILING DATE: 1998-06-26

PRIOR APPLICATION NUMBER: US 08/851,130
PRIOR FILING DATE: 1997-05-05

NUMBER OF SEQ ID NOS: 72

SEQ ID NO 72

LENGTH: L0

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APPLICANT: Sahin, Ugur
APPLICANT: Ramensee, Hans Georg
APPLICANT: Stevanovic, Stefan
APPLICANT: Stevanovic, Stefan
TITLE OF INVENTION: Method for Determining Presence of Cancer In a Sample By Determin
TITLE OF INVENTION: Gene, and Uses Thereof
TITLE OF INVENTION WIMBER: US/09/344,040C
CURRENT FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: US 09/105,839
PRIOR APPLICATION NUMBER: US 09/105,839
PRIOR APPLICATION NUMBER: US 08/851,130
PRIOR FILING DATE: 1997-05-05
NUMBER OF SEQ ID NOS: 132
SEQ ID NO 114
LENGTH: 10
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                                                                                                                                                                                                                                                                                       Query Match 53.8%; Score 21; DB 1; Length 10; Best Local Similarity 100.0%; Pred. No. 2.6e+02; Matches 4; Conservative 0; Mismatches 0; Indels
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; LENGTH: 10 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-397-633A-93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 50.0
Matches 3; Conservative
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CRGANISM: Homo sapiens
US-09-105-839D-72
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                                                               Query Match 53.8%; Score 21; DB 4; Length 10; Best Local Similarity 50.0%; Pred. No. 2.6e+02; Matches 3; Conservative 2; Mismatches 1; Indels
; ORGANISM: Homo sapiens US-09-344-040C-114
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GenCore version 5.1.6
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- protein search, using sw model OM protein

Run on:

October 21, 2004, 07:12:54; Search time 15.75 Seconds (without alignments) 54.981 Million cell updates/sec

US-09-720-469A-37 39 1 KXHRVIKDX 9

Title: Perfect score: Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

283416 seqs, 96216763 residues Searched:

1582

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 12

Post-processing: Minimum Match 00%
Maximum Match 100%
Listing first 45 summaries

PIR Database

1: piri: * 2: piri: * 3: piri: * 4: pir4: * **79:**

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	c and t	glycine reductase	e,	Ig heavy chain CDR	omeric	Ig heavy chain CRD	ribosomal protein	serum heterodimer,	unidentified 6.0/1	О	alcohol dehydrogen	collagen alpha 1(I	T-cell receptor be	cytochrome-c oxida	ferredoxin a2 - Ja	hypothetical prote	•-	macrophage cytotox	no-acid ox	virG protein - Agr	cytochrome-c oxida	neurokinin A - rai	neurokinin A - Atl	T-cell receptor al	Ig gamma-2b chain	alpha-myosin heavy	dihydrolipoamide S	parathyroid hormon	glycine cleavage s
SUMMARIES		C4374	B39308	152708	PT0228	PC4377	44	S78420	B56899	PQ0733	S09082	802617	B56979	PT0520	865381	869165	T31612	A42266	A60427	S39437	S13224	T14212	S23307	S23186	S23364	C30503	I46868	B45796		25
	DB	7	2	4	7	7	7	7	7	N	~	N	7	~	7	7	7	N	7	7	7	7	7	~	7	7	7	7	7	7
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	Result No.	! !	2	c	4	2	9	7	c o	on.	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25		27	28	29

N-methylpurine DNA	peptide-N4- (N-acet	cytochrome-c oxida	bothropstoxin - ja	pyruvate synthase	ribosomal protein	DNA-binding protei	hemoglobin, extrac	NAD ADP-ribosyltra	unidentified 5.4/3	hydrogensulfite re	hypothetical prote	triacylglycerol li	Ig heavy chain CRD	T-cell receptor al	gene NF2 protein -
A56029	JN0440	877980	A30823	H54346	S78765	C34135	S65730	S21163	PQ0730	S11024	B44510	S57274	PT0298	A35768	154379
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C;Species: Rattus norvegicus (Norway rat)
C;Spate: 28-Oct-1997 #sequence_revision 28-Oct-1997 #text_change 07-May-1999
C;Accession: PC4374
R;Sarig, G.; Weisman-Shomer, P.; Fry, M.
Biochem. Biophys. Res. Commun. 237, 617-623, 1997
A;Title: Telomeric and tetraplex DNA binding properties of qTBP42: A homologue of the CA1A;Reference number: PC4371; MUID:97445086; PMID:9299414
A;Accession: PC4374
A;Molecule type: protein
A;Residues: 1-10 <AR>
C;Comment: This protein binds either strand of the telomeric DNA as well as unimolecular
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Matches 3; Conserv
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6 KEHRL 10 1 KXHRV 5 8 d

RESULT 2 B39308 glycine reductase (EC 1.4.99) sulfhydryl protein C, beta chain - Clostridium sticklandi C;Species: Clostridium sticklandii	revision 19-Jun-1992 #text_change 09-Jul-2004	0. biol. Chem. 288, 2214/-22153, 1991 A;Title: Glycine reductase protein C. Properties and characterization of its role in the A;Reference number: A39308; MUID:92042141; PMID:1939235			EV92	A;Description: glycine reductase complex catalyzes the reductive deamination of glycine t C;Keywords: ATP; oxidoreductase	41.0%; Score 16; DB 2; Length 10; 50.0%; Pred. No. 2.5e+03;	2; Mismatches 1; Indels 0; Gaps 0;
RESULT 2 B39308 glycine reductase (EC 1.4.99) sulfhydryl protein C, beta C:Species: Clostridium sticklandii	C;Date: 19-Jun-1992 #sequence_revision 19-Jun-1992 #text_change 09-Jul-2004 C;Accession: B39308 R;Stadtman, T.C.; Davis, J.N.	U. BIOI. CHell. 200, 2214/-22153, 1991 A;Title: Glycine reductase protein C. Properties and chara. A;Reference number: A39308; MUID:92042141; PMID:1939235	A;Accession: B39308 A;Status: preliminary	A; Molecule type: protein A; Residues: 1-10 <sta></sta>	A;Cross-references: UNIPROT:Q9EV92 C;Function:	A;Description: glycine reductase complex catalyzes the rec C;Keywords: ATP; oxidoreductase	Query Match 41.0%; Score 16; DB 2; Lengt? Best Local Similarity 50.0%; Pred. No. 2.5e+03;	ive

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C; Accession: PT0246
R; Yamada, M.; Wassexman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
A; Title: Preferential utilization of specific immunoglobulin heavy chain diversity and jc A; Reference number: PT0222; MUID:91108337; PMID:1899102
A; Accession: PT0246
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C; Date: 11-Aug-1995 #sequence_revision 11-Aug-1995 #text_change 09-Jul-2004
C; Accession: B56899
R; Vazquez-Moreno, L.; Forath, J.; Schluter, S.F.; Marchalonis, J.J.
Comp. Biochem. Physiol. B 103, 563-568, 1992
Comp. Biochem. Physiol. B 103, 563-568, 1992
A; Title: Purification of a novel heterodimer from shark (Carcharhinus plumbeus) serum by A; Reference number: A56899; MUID:93092592; PMID:1458832
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PT0246
Ig heavy chain CRD3 region (clone 2-103D) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; Species: Rattus norvegicus (Norway rat)
C; Date: 25-Feb-1998 #sequence_revision 13-Mar-1998 #text_change 21-Jul-2000
C; Accession: S78420
R; Goldschmidt-Reisin, S.; Graack, H.R.
submitted to the Protein Sequence Database, February 1998
A; Reference number: S78411
A; Accession: S78420
                                                                                                    Gaps
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A;Residues: 1-9 <GOL>
A;Note: the protein is designated as mitochondrial ribosomal protein L41
C;Keywords: mitochondrion; protein biosynthesis; ribosome
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Length 12;
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Score 15; DB 2; I
Pred. No. 4.9e+03;
0; Mismatches 2;
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Best Local Similarity 66.7%; Pred. No. 2.8e+05;
Matches 2; Conservative 1; Mismatches 0.
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A;Molecule type: protein
A;Residues: 1-10 <VAZ>
A;Cross-references: UNIPROT:Q7LZJ7
   38.5%;
60.0%;
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                                              Similarity 60.0
3; Conservative
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Best Local Similarity
Matches 2; Conserv
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A; Residues: 1-7 < YAM>
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Query Match
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Matches 3
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S78420
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                                                                                ELECTION To the neuronal protein 1, truncated splice form - human
N;Alternate names: Drosophila ELAV(embryonic lethal, abnormal vision)-like 4; Hu antigen
C;Species: Homo sapiens (man)
C;Date: 18-Aug-2000 #sequence_revision 18-Aug-2000 #text_change 09-Jul-2004
C;Accession: 152708
R;Sekido, Y.; Bader, S.A.; Carbone, D.P.; Johnson, B.E.; Minna, J.D.
Cancer Res. 54, 4988-4992, 1994
A;Title: Molecular analysis of the HuD gene encoding a paraneoplastic encephalomyelitis
A;Reference number: 152708; MUID:94349312; PMID:8069866
A;Accession: 152708
A;Accession: L52708
A;Accession: L52708
A;Accession: L52708
A;Accession: L52708
A;Accession: L52708
A;Residues: 1-11 < SEMENA
A;Residues: 1-11 < SEM
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A,Residues: 1-12 <SAR>
A,Cross-references: UNIPROT:Q9QX80; UNIPROT:Q9Z0U8; UNIPROT:088311; UNIPROT:Q9QX81
C;Comment: This protein binds either strand of the telomeric DNA as well as unimolecular
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PT0228
Ig heavy chain CDR3 region (clone 1-112) - human (fragment)
C; Species: Homo sapiens (man)
C; Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C; Accession: PT0228
R; Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.J. Exp. Med. 173, 395-407, 1991
A; Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A; Reference number: PT0222; MUID:91108337; PMID:1899102
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 28-Oct-1997 #sequence_revision 28-Oct-1997 #text_change 09-Jul-2004
C;Accession: PC4377
R;Sarig, G.; Weisman-Shomer, P.; Fry, M.
Biochem. Biophys. Res. Commun. 237, 617-623, 1997
A;Title: Telomeric and tetraplex DNA binding properties of qTBP42: A homologue of the (A;Reference number: PC4371; MUID:97445086; PMID:9299414
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A;Residues: 1-12 <YAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
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Best Local Similarity 40.00
The 2; Conservative
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nes 2; Conservative
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RYVRD 6
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RILK 9
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PC4377
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Query Match Best Local S

Matches

RESULT 9

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N;Alternate names: collagen alpha 3(XI) chain
C;Species: Bos primigenius taurus (cattle)
C;Date: 03-Oct-1995 #sequence_revision 03-Oct-1995 #text_change 03-Oct-1995
C;Accession: B56979
R;Wu, J.J.; Eyre, D.R.
J. Biol. Chem. 270, 18865-18870, 1995
A;Title: Structural analysis of cross-linking domains in cartilage type XI collagen. Insi
A;Reference number: A56978; MUID:95370194; PMID:7642541
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-6 <WUA>
A;Note: the residue designated "X" is modified Lysine in collagen 1(II) some cross-linked
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A; Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A; Reference number: PT0509; MUID:91277601; PMID:1711558
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A, Cross-references: UNIPROT: P19854
C, Superfamily: alcohol dehydrogenase; long-chain alcohol dehydrogenase homology
C, Keywords: alcohol metabolism; NAD; oxidoreductase
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0520
R;Feeney, A.J.
U. Exp. Med. 174, 115-124, 1991
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                                                                                                        33.3%; Score 13; DB 2; Length 6; 100.0%; Pred. No. 2.8e+05; vative 0; Mismatches 0; Indels
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Pred. No. 2.8e+05;
0; Mismatches 1;
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A;Status: translation not shown
A;Molecule type: mRNA
A;Residues: 1-7 <FEE>
A;Experimental source: adult thymus, strain BALB/c
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                          collagen alpha 1(II) chain - bovine (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         33.3%;
66.7%;
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S65381
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C;Species: Equus caballus (domestic horse)
C;Species: Equus caballus (domestic horse)
C;Date: 28-Feb-1990 #sequence_revision 28-Feb-1990 #text_change 09-Jul-2004
C;Accession: S02617
R;Fairwell, T.; Julia, P.; Kaiser, R.; Holmquist, B.; Pares, X.; Vallee, B.L.; Joernvall FEBS Lett. 222, 99-103, 1987
A;Fitle: Acctylated N-terminal structures of class III alcohol dehydrogenases. Difference A;Reference number: S02617; MUID:88005160; PMID:3653405
A;Accession: S02617
A;Molecule type: protein
A;Residues: 1, 6 < FAI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Droteasome chain 1 - rat (fragment)

N;Alternate names: multicatalytic proteinase chain 1

C;Species: Rattus norvegicus (Norway rat)

C;Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change 21-Aug-1998

C;Accession: 809082

R;Lilley, K.S.; Davison, M.D.; Rivett, A.J.

FEBS Lett. 262, 327-329, 1990

A;Title: N-terminal sequence similarities between components of the multicatalytic prote
A;Reference number: 809082; MUID:90242957; PMID:2335214
                                                                                                                                                                                                                                                                                                                                             unidentified 6.0/15K protein [imported] - rice (fragment)
C;Species: Oryza sativa (rice)
C;Species: Oryza sativa (rice)
C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 20-Apr-2000
C;Accession: PQ0733
R;Komatsu, S.; Kajiwara, H.; Hirano, H.
Theor. Appl. Genet. 86, 935-942, 1993
A;Title: A rice protein library; a data-file of rice proteins separated by two-dimension A;Reference number: PQ0696
A;Accession: PQ0733
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                                                    35.9%; Score 14; DB 2; Length 10; ilarity 40.0%; Pred. No. 6.7e+03; Conservative 2; Mismatches 1; Indels
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A,Residues: 1-12 <LIL>
C,Superfamily: multicatalytic endopeptidase complex chain C9
    C;Keywords: glycoprotein; plasma
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Best Local Similarity 50.0
Matches 2; Conservative
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A, Molecule type: protein
A, Residues: 1-11 < KOM>
                                                                       Local Similarity
ses 2; Conserv
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6 RVVNE 10
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RESULT 10

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Query Match Best Loc Matches us-09-720-469a-37.rpr

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A;Title: Cytochrome-c oxidase in developing rat heart. Enzymic properties and amino-term A;Reference number: S65372; MUID:95324529; PMID:7601105
A;Accession: S65381
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-8 <SCH>
A;Cross-references: UNIPROT:P80430
C;Keywords: oxidoreductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 15
S69165
ferredoxin a2 - Japanese radish (fragment)
C;Species: Kaiware daikon (Japanese radish)
C;Species: Raiware daikon (Japanese radish)
C;Date: 10-Mar-1998 #sequence_revision 10-Mar-1998 #text_change 09-Jul-2004
C;Accession: S69165
R;Obata, S:; Nishimura, M.; Nagai, K.; Sakihama, N.; Shin, M.
Arch. Biochem. Biophys. 316, 797-802, 1995
A;Title: Four ferredoxins from Japanese radish leaves.
A;Title: Four ferredoxins from Japanese radish leaves.
A;Reference number: S69164; MUID:95168867; PMID:7864635
A;Molecule type: protein
A;Residues: 1-8 <0BA>
A;Cross-references: UNIPROT:Q7MIF1
C;Keywords: 2Fe-2S; electron transfer; iron-sulfur protein
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                                                                                                                                                                                                                                                                                                                               0; Indels
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Search completed: October 21, 2004, 07:27:57 Job time : 17.75 secs

3 HR 4

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2 HR 3

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Sequence 1, Application US/10788016
Publication No. US20040141992A1
GENERAL INFORMATION:
APPLICANT: ITOH, Kyogo
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US-10-447-161-84
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US-10-788-016-1
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                                                                                                                                                            ; Search time 61.75 Seconds (without alignments) 47.187 Million cell updates/sec
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| cgn2 6/ptodata/1/pubpaa/USO7_PUBCOMB.pep:*
| cgn2 6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
| cgn2 6/ptodata/1/pubpaa/USO6_PUBCOMB.pep:*
| cgn2 6/ptodata/1/pubpaa/USO6_PUBCOMB.pep:*
| cgn2 6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
| cgn2 6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*
| cgn2 6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*
| cgn2 6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*
| cgn2 6/ptodata/1/pubpaa/USO9B_PUBCOMB.pep:*
| cgn2 6/ptodata/1/pubpaa/USO0B_PUBCOMB.pep:*
| cgn2 6/ptodata/1/pubpaa/USO0B_PUBCOMB.pep:*
| cgn2 6/ptodata/1/pubpaa/USO0B_PUBCOMB.pep:*
| cgn2 6/ptodata/1/pubpaa/USOOB_PUBCOMB.pep:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-10-788-016-1

US-09-984-056-61

US-09-984-056-62

US-09-984-057-61

US-09-984-057-62

US-10-105-232-61

US-10-105-232-62

US-10-105-232-62

US-10-105-232-62

US-10-105-232-62

US-10-11-207-23

US-10-211-207-23

US-10-211-207-23

US-10-211-207-23

US-10-077-106-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1364641 seqs, 323758627 residues
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                                                                                                          - protein search, using sw model
                                                                                                                                                               October 21, 2004, 07:15:25
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Gapop 10.0 , Gapext 0.5
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Match Length
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Perfect score:
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Sequence 14, Appl
Sequence 114, Appl
Sequence 12, Appl
Sequence 114, Appl
Sequence 173, Appl
Sequence 578, Appl
Sequence 56, Appl
Sequence 54, Appl
Sequence 314, Appl
Sequence 314, Appl
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Sequence 314, Appl
Sequence 48, Appl
Sequence 54, Appl
          Sequence 72, Appl
Sequence 72, Appl
Sequence 72, Appl
Sequence 72, Appl
Sequence 145, Appl
Sequence 14, Appl
Sequence 22, Appl
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US-09-851-138-204

US-09-984-056-72

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US-10-189-437-72

US-10-293-822-14

US-10-293-822-14

US-10-293-822-22

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US-09-876-904A-578

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US-09-305-982-46

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ALIGNMENTS

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Gaps
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0
                                                                   APPLICANT: Wang, Rong-fu
TITLE OF INVENTION: Mutant Fibronectin and Tumor Metastasis
FILE REFERENCE: HO-P02484US1
CURRENT APPLICATION NUMBER: US/10/447,161
CURRENT APPLICATION NUMBER: 60/383,530
PRIOR APPLICATION NUMBER: 60/383,530
PRIOR FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 148
SOFTWARE: Patentin version 3.1
SEQ ID NO 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92.3%; Score 36; DB 15; Length 9; 87.5%; Pred. No. 1.2e+06; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Synthetic Peptide
Sequence 84, Application US/10447161 Publication No. US20040023314A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Artificial Sequence
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TITLE OF INVENTION: ANTHRAX AND SMALL POX REPLIKINS AND METHODS OF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: BOGOCH, ELENORE S.
TITLE OF INVENTION: REPLIKINS AND METHODS OF IDENTIFYING
TITLE OF INVENTION: REPLIKIN-CONTAINING SEQUENCES
FILE REFERENCE: 09425-46902
CURRENT APPLICATION NUMBER: US/09/984,057
CURRENT FILING DATE: 2001-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 28; DB 9; I
Pred. No. 1.2e+06;
1; Mismatches 2;
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62.5%; Pred. No. 1.2e+06;
cive 1; Mismatches 2
                    FILE REFERENCE: 09425-46903;
CURRENT APPLICATION NUMBER: US/09/984,056
CURRENT FILING DATE: 2001-10-26
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR FILING DATE: 2001-07-09
PRIOR PILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR APPLICATION NUMBER: 09/146,755
PRIOR FILING DATE: 1998-09-04
PRIOR FILING DATE: 1998-09-04
PRIOR FILING DATE: 1998-09-04
PRIOR FILING DATE: 1994-02-17
PRIOR FILING DATE: 1994-02-17
PRIOR FILING DATE: 1994-02-17
PRIOR FILING DATE: 1994-02-17
PRIOR PELING DATE: 1994-02-17
PRIOR PELING DATE: 1994-02-17
PRIOR APPLICATION NUMBER: 08/198,139
PRIOR FILING DATE: 1994-02-17
PRIOR FILING DATE: 1994-02-17
PRIOR PRIOR PRIOR NOS: 103
PRIOR FILING DATE: 1994-02-17
PRIOR PRIOR PRIOR NOS: 103
PRIOR FILING DATE: 1994-02-17
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PRIOR APPLICATION NUMBER: 60/278,761
PRIOR FILING DATE: 2001-07-09
PRIOR FILING DATE: 2001-03-27
PRIOR FILING DATE: 2001-03-27
PRIOR FILING DATE: 1998-09-04
PRIOR FILING DATE: 1998-09-04
PRIOR FILING DATE: 1998-09-04
PRIOR FILING DATE: 1994-02-17
PRIOR FILING DATE: 1994-02-17
PRIOR FILING DATE: 1994-02-17
NUMBER OF SEQ ID NOS: 90
SOFTWARE: Patentin 2.1
SEQ ID NO 61
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US-09-984-057-62
; Sequence 62, Application US/09984057
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Patent No. US20020151677A1
GENERAL INFORMATION:
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62.5%;
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Best Local Similarity 62.5
Matches 5; Conservative
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Matches 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: Polyoma virus
US-09-984-057-61
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US-09-984-056-62
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US-09-984-057-61
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                                                                                                                                                                                                                                                                                                                                                                               CTHER INFORMATION: Peptide consisting of 9 amino acid residues from the 84th residue;
CTHER INFORMATION: to the 92nd residue of cyclophillin B
US-10-788-016-1
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US-09-984-056-61

i Sequence 61, Application US/09984056

j Patent No. US2020120106A1

j GENERAL INFORMATION:

APPLICANT: BOGOCH, ELENORE S.

TITLE OF INVENTION: ANTHRAX AND SMALL POX REPLIKINS AND METHODS OF USE

FILE REPERENCE: 09425-46903

CURRENT FILING DATE: 2001-10-26

PRIOR APPLICATION NUMBER: 60/303,396

PRIOR APPLICATION NUMBER: 60/278,761

PRIOR APPLICATION NUMBER: 09/146,755

PRIOR FILING DATE: 2001-03-27

PRIOR PLILNG DATE: 2001-03-27

PRIOR PLILNG DATE: 1998-09-04

PRIOR PLILNG DATE: 1998-09-04

PRIOR FLILNG DATE: 1994-02-17

NUMBER OF SEQ ID NOS: 103

SOFTWARE: Patentin 2.1
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                            FILE REFERENCE: 3190-049

CURRENT APPLICATION NUMBER: US/10/788,016

CURRENT FILING DATE: 2004-02-26

PRIOR APPLICATION NUMBER: PCT/JP02/08641

PRIOR FILING DATE: 2002-08-28

PRIOR FILING DATE: 2001-08-29

PRIOR FILING DATE: 2001-08-29

NUMBER OF SEQ ID NOS: 9

SOFTWARE: Patentin version 3.2

; SEQ ID NO 1

LENGTH: 9
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            TITLE OF INVENTION: Desensitizers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92.3%;
87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 62.5
Matches 5; Conservative
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KQHRELKD 8
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                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: Artificial
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US-09-984-056-62
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US20020151677A1

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APPLICANT: BOGOCH, SAMUEL
APPLICANT: BOGOCH, ELENORE S.
TITLE OF INVENTION: REPLIKIN PEPTIDES IN RAPID REPLICATION OF GLIOMA CELLS
TITLE OF INVENTION: AND IN INFLUENZA EPIDEMICS
FILE REFERENCE: 09425-46904
CURRENT APPLICATION NUMBER: US/10/105,232
CURRENT FILING DATE: 2002-03-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: BOGOCH, SAMUEL
APPLICANT: BOGOCH, ELENORE S.
TITLE OF INVENTION: REPLIKIN PEPTIDES AND ANTIBODIES THEREFORE
FILE REFERENCE: 09425/46905
CURRENT APPLICATION NUMBER: US/10/189,437
CURRENT FILING DATE: 2002-07-08
PRIOR APPLICATION NUMBER: 10/105,232
PRIOR FILING DATE: 2002-03-26
PRIOR FILING DATE: 2001-10-26
PRIOR PPLICATION NUMBER: 60/303,396
PRIOR FILING DATE: 2001-07-09
PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 729
SOFTWARE: PATENTIN 2.1
SEQ ID NO 61
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         71.8%; Score 28; DB 14; Length 9; 62.5%; Pred. No. 1.2e+06; ative 1; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/303,396
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR PILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR APPLICATION NUMBER: 09/146,755
PRIOR APPLICATION NUMBER: 09/817,144
PRIOR APPLICATION NUMBER: 09/817,144
PRIOR FILING DATE: 2001-03-27
PRIOR FILING DATE: 2001-03-27
PRIOR PILING DATE: 1994-02-17
NUMBER OF SEQ ID NOS: 535
SOFTWARE: Patentin 2.1
SEQ ID NO 62
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                                                                    US-10-105-232-62; Application US/10105232; Publication No. US20030180328A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5; Conservative
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US-10-189-437-61
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Best Local Similarity
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1 KQHRELKD
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APPLICANT: BOGOCH, SAUGEL

APPLICANT: BOGOCH, ELENORE S.

TITLE OF INVENTION: REPLIKIN PEPTIDES IN RAPID REPLICATION OF GLIOMA CELLS

TITLE OF INVENTION: AND IN INFLUENZA EPIDEMICS

TITLE OF INVENTION: AND IN INFLUENZA EPIDEMICS

FILE REFERENCE: 09425-46904

CURRENT APPLICATION NUMBER: US/10/105,232

CURRENT FILING DATE: 2002-03-26

PRIOR APPLICATION NUMBER: 60/278,761

PRIOR FILING DATE: 2001-03-27

PRIOR FILING DATE: 1998-09-04

PRIOR FILING DATE: 1998-09-04

PRIOR FILING DATE: 1994-02-17

NUMBER: O8/198,139

PRIOR FILING DATE: 1994-02-17

NUMBER: PALCATION NUMBER: 08/198,139

PRIOR FILING DATE: 2001-03-27

PRIOR FILING DATE: 2001-03-27

PRIOR FILING DATE: 2001-03-27

PRIOR FILING DATE: 2001-03-27

PRIOR FILING DATE: 1994-02-17

NUMBER: PALCATION NUMBER: 08/198,139

PRIOR FILING DATE: 1904-02-17

SEQ ID NOS: 535

SEQ ID NO 61
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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: BOGOCH, SAMUEL

APPLICANT: BOGOCH, SAMUEL

APPLICANT: BOGOCH, SAMUEL

TITLE OF INVENTION: REPLIKIN-CONTAINING SEQUENCES

TITLE OF INVENTION: REPLIKIN-CONTAINING SEQUENCES

FILE REFERENCE: 09425-46902

CURRENT APPLICATION NUMBER: US/09/984,057

CURRENT FILING DATE: 2001-026

PRIOR FILING DATE: 2001-07-09

PRIOR FILING DATE: 2001-07-09

PRIOR FILING DATE: 2001-03-27

PRIOR FILING DATE: 1998-09-04

PRIOR FILING DATE: 1998-09-04

PRIOR FILING DATE: 1998-03-07

SOFTWARE: PALENTING DATE: 1994-02-17

NUMBER OF SEQ ID NOS: 90

SSEQ ID NO 62
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Pred. No. 1.2e+06;
1; Mismatches 2; Indels
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Publication No. US20030180328A1
GENERAL INFORMATION:
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62.5%;
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Best Local Similarity 62.5
Matches 5; Conservative
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CORGANISM: Polyoma virus
US-09-984-057-62
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; ORGANISM: Polyoma virus
US-10-105-232-61
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Best Local Similarity
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APPLICANT: COLLING, MALY
APPLICANT: Madrenas, Joaquin
APPLICANT: Madrenas, Deatriz
APPLICANT: Carreno, Beatriz
APPLICANT: Kuchroo, Vijay
TITLE OF INVENTION: METHODS FOR MODULATING AN IMMUNE
TITLE OF INVENTION: PP2A
FILE REFERENCE: GNN-027
CURRENT APPLICATION NUMBER: US/10/211,207
CURRENT APPLICATION NUMBER: US/10/077,106
PRIOR APPLICATION NUMBER: US/10/077,106
PRIOR FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 24
LENGTH: 9
                                                            Sequence 24, Application US/10211207
Publication No. US20030004113A1
GENERAL INFORMATION:
APPLICANT: Collins, Mary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 66.7
Matches 4; Conservative
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Best Local Similarity
Matches 4; Conserv
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HRELKO
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US-10-077-106-24
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US-10-077-106-23
                                           US-10-211-207-24
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Sequence 23, Application US/10211207

Bublication No. US2030004113A1

GENERAL INFORMATION:

APPLICANT: Collins, Mary

APPLICANT: Collins, Mary

APPLICANT: Carreno, Beatriz

APPLICANT: Carreno, Wijay

TITLE OF INVENTION: RESPONSE BY MODULATING AN IMMUNE

TITLE OF INVENTION: PP2A

TITLE OF INVENTION: PP2A

TITLE OF INVENTION: PP2A

TITLE OF INVENTION: PP2A

TITLE OF SINCE SONO NUMBER: US/10/211,207

CURRENT FILING DATE: 2002-08-02

PRIOR APPLICATION NUMBER: US/10/077,106

PRIOR APPLICATION NUMBER: US/10/077,106

PRIOR APPLICATION NUMBER: US/10/077,106

NUMBER OF SEQ ID NOS: 25

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 23

LENGTH: P
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                                                   RESULT 10
US-10-189-437-62

Sequence 62, Application US/10189437

Publication No. US20030194414A1

GENERAL INFORMATION:

APPLICANT: BOGOCH, SAMUEL

APPLICANT: BOGOCH, ELENORE S.

TITLE OF INVENTION: REPLIKIN PEPTIDES AND ANTIBODIES THEREFORE

FILE REFERENCE: 09425/46905

CURRENT APPLICATION NUMBER: US/10/189,437

CURRENT FILING DATE: 2002-07-08

PRIOR FILING DATE: 2002-03-26

PRIOR FILING DATE: 2001-10-26

PRIOR FILING DATE: 2001-10-26

PRIOR PULICATION NUMBER: 60/303,396

PRIOR FILING DATE: 2001-07-09

PRIOR FILING DATE: 2001-03-27

NUMBER OF SEQ ID NOS: 729

SOFTWARE: Patentin 2:1

SEQ ID NO 62

LENGTH: 9

TAVED: DATE
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Best Local Similarity 62.5
Matches 5; Conservative
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Best Local Similarity
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HRELKD 6
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TYPE: PRT
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APPLICANT: Collins, Mary
APPLICANT: Collins, Mary
APPLICANT: Carreno, Beatriz
APPLICANT: Carreno, Vijay
TITLE OF INVENTION: METHODS FOR MODULATING THE INTERACTION BETWEEN CTLA4 AND
TITLE OF INVENTION: PP2A
FILLE OF INVENT
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                                                                                                                                                                                                                                                 61.5%; Score 24; DB 14; Length 9; 66.7%; Pred. No. 1.2e+06; cive 1; Mismatches 1; Indels
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; Publication No. US20030166531A1
; GENERAL INFORMATION:
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TYPE: PRT
CRGANISM: Polyomavirus medium
US-10-211-207-24
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COGANISM: Polyomavirus small
US-10-077-106-23
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STUYVER, LIEVEN
TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES
AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC
AGENTS
GENERAL INFORMATION:

APPLICANT: Collins, Mary
APPLICANT: Carreno, Beatriz
APPLICANT: Carreno, Wijay
TITLE OF INVENTION: METHODS FOR MODULATING THE INTERACTION BETWEEN CTLA4 AND
TITLE OF INVENTION: RESPONSE BY MODULATING THE INTERACTION BETWEEN CTLA4 AND
TITLE OF INVENTION: RESPONSE BY MODULATING THE INTERACTION BETWEEN CURRENT APPLICATION NUMBER: US/10/077,106
CURRENT APPLICATION NUMBER: 60/269,757
PRIOR APPLICATION NUMBER: 60/269,757
PRIOR FILING DATE: 2001-02-16
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FastSEQ for Windows Version 4.0
SOFTWARE: FastSEQ for Windows Version 4.0
SOFTWARE: PASTSEQ FOR WINDOWS VERSION 4.0
SOFTWARE: PASTSEQ FOR WINDOWS VERSION 4.0
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COUNTER: USAS

ZIP: 77210-4433

ZIP: 77220-4433

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Microsoft Word 6.0 / ASCII text output

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/851,138

FILING DATE: 09-May-2001

PRIOR APPLICATION NUMBER: EP 94870166.9

FILING DATE: 20 Jun 1995

ATTORNEY/AGENT INFORMATION:

NAME: KAMMERER, PATRICIA A.

REGISTRATION NUMBER: 29,775

REFERENCE/DOCKET NUMBER: 1NNS:004

INFORMATION FOR SEQ ID NO: 204:

SEQUENCE CHARACTERISTICS:

CHARACTERISTICS:

COMPUTER: AND AND ACTOR A.

REFERENCE/DOCKET NUMBER: 1NNS:004

INFORMATION FOR SEQ ID NO: 204:

SEQUENCE CHARACTERISTICS:

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COMPUTER: ADDITION ADDITION ADDITION ADDITION FOR SEQ ID NO: 204:

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ADDRESSEE: ARNOLD, WHITE & DURKEE STREET: P.O. BOX 4433
CITY: HOUSTON
STATE: TEXAS
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Best Local Similarity 66.7%;
Matches 4; Conservative
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TOPOLOGY: linear
MOLECULE TYPE: peptide
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HRELKD 6
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SEQUENCE DESCRIPTION: SEQ ID NO: 204:
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US-09-851-138-204
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Search completed: October 21, 2004, 07:33:41 Job time: 62.75 secs

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Cygnus columbianus (tundra swan).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Anseriformes; Anatidae; Cygnus.
NCBI_TaxID=110926;
                                                                                                                                                                                                                      Anas platyrhynchos (Domestic duck).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleos
Archosauria; Aves; Neognathae; Anseriformes; Anatidae; Anas.
NCBI_TaxID=8839;
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                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
Cotter J.-P., StJohn J., Quinn T.W.;
Submitted (JUN-2002) to the EMBL/GenBank/DDBJ databases
EMBL; AY123327; AAM96901.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
Cotter J.-P., StJohn J., Quinn T.W.;
Submitted (JUN-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AY123326; AAM96900.1; -.
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LDH-B (Fragment).
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                                                     October 21, 2004, 07:05:04; Search time 86 Seconds (without alignments) 60.214 Million cell updates/sec
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Copyright (c) 1993 - 2004 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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DB 2; Length 8;

Euteleostomi;

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01-MAR-2004 (TrEMBLrel. 26, 01-MAR-2004 (TrEMBLrel. 26, 01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 60.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRELIMINARY;
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Matches 3; Conserv
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                                                                                                                                                                                                                                                                               NCBI_TaxID=4932;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Anseriformes; Anatidae; Coscoroba.
[13] TaxID=8863;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anser caerulescens caerulescens (snow goose).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Anseriformes; Anatidae; Anser.
NCBI_TaxID=70340;
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Pred. No. 1.8e+06; .....hes 1; Indels
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Cotter J.-P., StJohn J., Quinn T.W.;
Submitted (JUN-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AY123324; AAM96898.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. Cotter J.-P., StJohn J., Quinn T.W.; Submitted (JUN-2002) to the EMBL/GenBank/DDBJ databases.
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Last annotation update)
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Last annotation update)
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                                                        3; Mismatches
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Best Local Similarity 33.5°,
Local 2; Conservative
                          Similarity 33.3
2; Conservative
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HKIVAD 7
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SEQUENCE
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SEQUENCE
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Q05403
ID Q0540
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                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                 Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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Sordariomycetidae, Sordariales, Sordariaceae, Neurospora.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -!- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which preliminary data.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 8;
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SEQUENCE 9 AA; 1173 MW; C483A32B50440449 CRC64;
                                                                       Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Last sequence update)
Last annotation update)
01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update
Orf 00915 protein (Fragment).
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60.0%; Pred. No. 1.8e+06;
iive 1; Mismatches 1;
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us-09-720-469a-37.rup

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PRT;
                                       48.78;
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 10 AA; 1206 MW;
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Best Local Similarity 60.v-
                                                                                   3; Conservative
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                                                                                                                                                                                                                                                                                PRELIMINARY;
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                                                                                                                                                                    4 RSHRPIR 10
                                                                                                                              1 KXHRVIK 7
                                                                Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                         Name=PDGFA;
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 SEQUENCE
                                           Query Match
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Matches
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Q9XSP2
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STRAIN=NC16-2, TC29-5, and TC97;
MEDLINE=22315381; PubMed=12427948;
Kholodii G., Gorlenko Z., Mindlin S., Hobman J., Nikiforov V.;
"Tn5041-like transposons: molecular diversity, evolutionary relationships and distribution of distinct variants in environmental
                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE=Late pregnancy myometrium;
MEDLINE=22181008; PubMed=12193412;
Yang M., Gupta A., Shlykov S.G., Corrigan B., Tsujimoto S.,
Sanborn B.M.;
"Multiple Trp isoforms implicated in capacitative calcium entry are expressed in human pregnant myometrium and myometrial cells.";
Biol. Reprod. 67:988-994(2002).
EMBL; AF483646; AAM97861.1; -.
                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
Pseudomonadaceae; Pseudomonas.
NCBI_TaxID=294;
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48.7%; Score 19; DB 2; Length 10;
Best Local Similarity 80.0%; Pred. No. 2.9e+03;
Matches 4; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ422128; CAD19527.1; -.
EMBL; AJ422129; CAD19528.1; -.
EMBL; AJ422130; CAD19529.1; -.
GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0005741; C:mitochondrial outer membrane; IEA.
GO; GO:0015288; F:porin activity; IEA.
                                                                                                                                                                                     01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
Capacitative calcium channel protein Trpl (Fragment).
Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10 AA; 1336 MW; 0DD0C0B401F40724 CRC64;
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01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Porin-like protein (Fragment).
                                                                                                                                                   10 AA
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STRAIN=NC16-2, TC29-5, and TC97;
Kholodii G.Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Microbiology 148:3569-3582(2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRELIMINARY;
                                                                                                                                                 PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pseudomonas fluorescens.
                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
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NRIYKD
HRVIKD
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QBRJF1;
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MEDLINE=20065871; PubMed=10598812;
Bonthron D.T., Smith S.L., Campbell R.;
"Complex patterns of intragenic polymorphism at the PDGFA locus.";
Hum. Genet. 105:452-459(1999).
EMBL; AJ243280; CAB45927.1; -.
NON_TER 1 1 1
NON_TER 11 11
SEQUENCE 11 AA; 1345 MW; 7FB881F101E1E044 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
MEDLINE=20065871; PubMed=10598812;
Bonthron D.T., Smith S.L., Campbell R.;
"Complex patterns of intragenic polymorphism at the PDGFA locus.";
Hum. Genet. 105:452-459(1999).
EMBL; AJ243277; CAB45926.1; -.
NON TER 1 1 1
NON TER 11 A3; 1331 MW; 7FB881F101E1F2D4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hylobates syndactylus (Siamang) (Symphalangus syndactylus).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hylobatidae, Hylobates.
NCBI_TaxID=9590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pan troglodytes (Chimpanzee).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Pan.
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                                                     Length 10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Indels
23C47E7401F5A417 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Platelet-derived growth factor A chain (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Platelet_derived growth factor A chain (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 19; DB 2; I
Pred. No. 3.2e+03;
1; Mismatches 1;
                                                   Score 19; DB 2; I Pred. No. 2.9e+03;
                                                                                                                                                                                                                                                                                                                                                     11 AA
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                                                                                                       2; Mismatches
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RESULT 11 Q9XSP6

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48.7%; Score 19; DB 2; I
60.0%; Pred. No. 3.2e+03;
tive 1; Mismatches 1;
 11 AA.
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Best Local Similarity 60..
"ne 3; Conservative
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PRELIMINARY;
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                                                                                                                                              NCBI_TaxID=98375;
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HRSVK 11
                                                                                            Presbytis johnii.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Name=PDGFA;
                                                                              Name=PDGFA;
                                                                                                                                    Presbytis.
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NON TER
SEQUENCE
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Q9XSP8
Q9XSP8;
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Q9XSQ4;
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Mammalia, Eutheria, Primates, Catarrhini, Cercopithecidae, Colobinae,
                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                        MEDLINE=20065871; PubMed=10598812;
Bonthron D.T., Smith S.L., Campbell R.;
"Complex patterns of intragenic polymorphism at the PDGFA locus.";
Hum. Genet. 105:452-459(1999).
EMBL; AJ243279; CAB45925.1; -.
NON TER 1 1 11
NON TER 11 11
SEQUENCE 11 AA; 1345 MW; 7FB881F101E1E044 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
MEDLINE=20065871; PubMed=10598812;
Bonthron D.T., Smith S.L., Campbell R.;
"Complex patterns of intragenic polymorphism at the PDGFA locus.";
Hum. Genet. 105:452-459(1999).
EMBL; AJ243282; CAB45924.1; -.
NON TER 1 1 1
NON TER 1 11
SEQUENCE 11 AA; 1345 MW; 7FB881F101B1E044 CRC64;
                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Pongo.
NCBI_TaxID=9600;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        48.7%; Score 19; DB 2; Length 11; 60.0%; Pred. No. 3.2e+03; ive 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                            48.7%; Score 19; DB 2; Length 11; 60.0%; Pred. No. 3.2e+03; ive 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Platelet-derived growth factor A chain (Fragment).
Name=PDGFA;
                                                                                                        01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Platelet-derived growth factor A chain (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pygathrix nemaeus (Dove langur)
                                                                                                                                                                           Pongo pygmaeus (Orangutan).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 60.00,
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Best Local Similarity 60..
". 3; Conservative
                                                                               PRELIMINARY;
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HRSVK 11
|| :|
HRSVK 11
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NCBI_TaxID=54133;
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                                                                                                                                                              Name=PDGFA;
              7
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                                                                                 94SX6D
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RESULT 12
Q9XSP7
ID Q9XSP
AC Q9XSP
AC Q9XSP
DT 01-NO
DT 01-DB
DT 01-DB
DE Plate
GN Name=
GN Name=
CC Mamma
OC Pygatl
OC Mamma
OC Pygatl
CC Mamma
OC Pygatl
CR Mamma
O

RESULT 13 Q9XSP8

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                                                                                                                                                                          Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Cercopithecidae, Colobinae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
MEDLINE=20065871; PubMed=10598812;
Bonthron D.T., Smith S.L., Campbell R.;
"Complex patterns of intragenic polymorphism at the PDGFA locus.";
Hum. Genet. 105:452-459(1999).
EMBL; AJ243281; CAB46013.1; -.
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MEDLINE=20065871; PubMed=10598812;
Bonthron D.T., Smith S.L., Campbell R.;
"Complex patterns of intragenic polymorphism at the PDGFA locus.";
Hum. Genet. 105:452-459(1999).
EMBL; AJ243278; CAB45916.1; -.
NON_TER 1 1 1
NON_TER 11 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gorilla gorilla (gorilla).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Gorilla.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Indels
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01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Platelet-derived growth factor A chain (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Platelet-derived growth factor A chain (Fragment).
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01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Hypothetical protein (Fragment).
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EXPLINED FROM N.A.

SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

STRAIN=17XNL;

R Carlton J.M., Angiuoli S.V., Suh B.B., Kooij T.W., Pertea M.,

A Carlton J.D., Pop M., Kosack D.S., Shumway M.F., Bidwell S.L.,

A Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Koo H.L.,

A Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Koo H.L.,

A Shallom S.J., van Aken S.E., Riedmuller S.B., Feldblyum T.V.,

A Cho J.K., Quackenbush J., Sedegah M., Shoaibi A., Cummings L.M.,

A Cho J.K., Quackenbush J., Sedegah M., Shoaibi A., Cummings L.M.,

A Cho J.K., Quackenbush J., Sedegah M., Shoaibi A., Cummings L.M.,

A Cho J.K., Quackenbush J., Sedegah M., Shoaibi A., Cummings L.M.,

A Van Lin L.H., Janse C.J., Waters R.R., Smith H.O., White O.R.,

A van Lin L.H., Janse C.J., Waters R.R., Smith H.O., White O.R.,

A van Lin L.H., Janse C.J., Waters R.R., Smith H.O., White O.R.,

A van Lin L.H., Janse C.J., Waters R.R., Smith H.O., White O.R.,

A van Lin L.H., Janse C.J., Waters R.R., Smith H.O., White O.R.,

A van Lin L.H., Janse C.J., Waters R.R., Smith H.O., White O.R.,

Carucci D.J.,

A carucci D.J.,

Carucci D.J.,

Carucci D.J.,

A carucci D.J.,

Carucci
               Plasmodium yoelii yoelii.
Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
NCBI_TaxID=73239;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 46.2%; Score 18; DB 2; Length 12; Best Local Similarity 60.0%; Pred. No. 5.8e+03; Matches 3; Conservative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12 12
12 AA; 1485 MW; D2A760BC2E633DC2 CRC64;
Name=PY04332;
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Search completed: October 21, 2004, 07:26:47 Job time : 87 secs

3 HRVIK 7 |:| | 3 HKVAK 7

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

October 21, 2004, 07:04:19; Search time 83 Seconds (without alignments) 38.898 Million cell updates/sec Run on:

US-09-720-469A-38 40 1 DXMIQGGDX 9 Title: Perfect score:

Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

2002273 seqs, 358729299 residues Searched:

497672 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 12

Post-processing: Minimum Match 08 Maximum Match 1008

Listing first 45 summaries

Database :

geneseqp1980s:*
geneseqp1990s:*
geneseqp2000s:*
geneseqp2002s:*
geneseqp2003as:*
geneseqp2003bs:* A_Geneseq_23Sep04:* 1: qeneseqp1980s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	769963 Human cyc	58 Human	Human	22 Human	5947 Human	Aab46931 Human cyc	89 Antitu	Abg79076 Human Cyp	74 Human			Adg89642 Class I H	_	_	Ade84662 Adenoviru	Aar20729 Pancreas-	Aay51905 Elastase	Aau78392 Biosurfac	Adk07318 Hepatitis	Adk06815 Hepatitis	Aaw83889 Peptide s	Aaw48850 Human pro	Aab45861 Human tum	3253	7045 Gi-
	ID Des	AAY69963 AAY6	AAY69958 Aay	61	922	7	н	AAG68089 Aag	ABG79076	4	ADC17709 Add	ADC17702 Add	ADG89642 Adç	AAY69957 Aay	AAB12089 Aal	N	AAR20729 Aa1	AAY51905 Aay	AAU78392 Aau	80	ADK06815 Ad	AAW83889 Aav	AAW48850 Aav	AAB45861 AaB	ABR83253 Abr8:	ADN17045 Adn1
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Gaps

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	Abr63060 Pathogen	Aae34120 Gluten sp	Aar93207 New contr	Aar72938 E. coli P	Aar72894 E. coli P	Abr67369 Human hsp	Abr67364 Human hsp	Adk07060 Hepatitis	Aar72949 E. coli P	Aar72905 E. coli P	Aaw41471 Fragment	Aab35991 Sorbitol	Aag87272 Saccharom	Adil5749 HLA-A2 re	Adj52136 CH1 delet	Adk09152 Human pap	Adk09635 Human pap	Adol7328 Different	Aar14821 Bay thioe
ABR01459	ABR63060	AAE34120	AAR93207	AAR72938	AAR72894	ABR67369	ABR67364	ADK07060	AAR72949	AAR72905	AAW41471	AAB35991	AAG87272	ADI15749	ADJ52136	ADK09152	ADK09635	ADO17328	AAR14821
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77	22	21	21	21	21	21	21	21	21	21	21	21	21	21	21	21	21	21	23
26	27	28				32	33		35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

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This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                 Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                         Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                92.5%; Score 37; DB 3; Length 9; ilarity 87.5%; Pred. No. 1.7e+06; Conservative 0; Mismatches 1; Indels
                                                                                         Human cyclophilin B peptide fragment #43
                                                                                                                                                                                                                                                                                                                                                                                                             Claim 10; Page 60; 64pp; Japanese.
              AAY69963 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                 (SUMU ) SUMITOMO PHARM CO LTD. (ITOH/) ITOH K.
                                                                                                                                                                                                                                   99WO-JP003360.
                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                              WPI; 2000-116932/10.
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Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                         Gomi S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9 AA;
                                                                                                                                                       Homo sapiens
                                                                                                                                                                                WO9967288-A1
                                                                                                                                                                                                                                  24-JUN-1999;
                                                                11-APR-2000
                                                                                                                                                                                                                                                         25-JUN-1998;
                                                                                                                                                                                                         29-DEC-1999,
                                       AAY69963;
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/label* Phe, Leu, Ile, Trp,
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                                                                                                    Gomi S;
                                                                                                                                                                                                              Sequence 9 AA;
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                WO9967288-A1
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                                              24-JUN-1999;
                                                             25-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                 This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                   Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                           Tumor antigen peptides derived from cyclophilin B for treatment and
                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                       1; Indels
                                                                                                                                                                                                                                                                                                                                                        Length 9;
                                                                                                                                                                                                                                                                                                                                                        Score 37; DB 3; I Pred. No. 1.7e+06;
                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /label= Phe, Tyr, Met, Trp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human cyclophilin B peptide fragment #41
                                                                                    Human cyclophilin B peptide fragment #38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY69961 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                    Claim 4; Page 61; 64pp; Japanese
                                       standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                        .0
                                                                                                                                                                                                       (SUMU ) SUMITOMO PHARM CO LID. (ITOH/) ITOH K.
                                                                                                                                                                                       98JP-00178449
                                                                                                                                                                                                                                                                                                                                                        92.5%;
                                                                                                                                                                         99WO-JP003360
                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
7; Conserve
                                                                                                                                                                                                                                                                     diagnosis of tumors.
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                                                                                                                                                                                                                                                                                                                                                                                      DXMIOGGD
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 DYMIQGGD
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                                                                                                                                                                                                                                                                                                                                         Sequence 9 AA;
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                                                                                                                                                                                        25-JUN-1998;
                                                                      11-APR-2000
                                                                                                                            Homo sapiens
                                                                                                                                                                         24-JUN-1999;
                                                                                                                                                         29-DEC-1999.
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                                        AAY69958
                                                       AAY69958;
                                                                                                                                                                                                                              Itoh K,
                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 3
                                AAY69961
ID AAX
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DT 11-
XXX
DE HUU
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XX
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CYC
KW HLA
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XX
CYC
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This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                     Tumor antigen peptides derived from cyclophilin E for treatment and diagnosis of tumors.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 92.5%; Score 37; DB 3; Le Best Local Similarity 100.0%; Pred. No. 1.7e+06; Matches 8; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human cyclophilin B peptide fragment #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 10; Page 60; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAY69922 standard; peptide; 9 AA.
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                                                                                                                     (SUMU ) SUMITOMO PHARM CO LID. (IIOH/) ITOH K.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99WO-JP003360.
                                                         98JP-00178449
99WO-JP003360
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us-09-720-469a-38.rag

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This invention describes the novel use of interferons (IFNS) or DNAS

capable of expressing the interferons and/or antigenic proteins (AP),

antigenic peptides derived from the proteins or DNAs capable of

expressing the antigeners or peptides, in the manufacture of an

agent for induction of antigen-specific T cells. The products of the

invention have virucide and cytostatic activity and can be used for gene

therapy or as inducers of antigen-specific T cells. The action of

therapy or as inducers of antigen-specific T cells. The action of

T cell (CTL) by administering an antigenic peptide in an incomplete

Freund's adjuvant (IFA) emulsion preparation form was evaluated. IFNS (or

DNA encoding IFNS) are useful in the manufacture of a medicament for

inducing antigen-specific T cells in an individual who has been

administered with AP (or DNA encoding AP) or vice versa. The medicament

is useful for the treatment or prophylaxis of a tumor or a viral
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA,
T
This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tumor antigenic peptide; interferon; IFN; antigen-specific T cell; virucide; cytostatic; gene therapy; IFN-alpha; cytotoxic T cell; viral infectious disease; cyclophilin B; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Use of antigenic proteins, peptides, interferon or their encoding the manufacture of an agent for the induction of antigen-specific
                                                                                                                                                                  .
                                                                                                                                                                                                                                                                                                                                                                                                                                     Human cyclophilin B derived tumor antigenic peptide SEQ ID 33.
                                                                                                                          Score 37; DB 3; Length 9;
Pred. No. 1.7e+06;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 20; 25pp; English.
                                                                                                                                                                                                                                                                                                                             AAB46947 standard; peptide; 9 AA.
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                                                                                                                            92.5%;
                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                7; Conservative
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Best Local Similarity
Matches 7; Conserv
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                                                                                          Sequence 9 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                AAB46947;
                                                                                                                                                                                                                                                                                        RESULT 5
  80 X C C C C
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This invention describes the novel use of interferons (IFNs) or DNAs capable of expressing the interferons and/or antigenic proteins (AP), antigenic peptides derived from the proteins or DNAs capable of expressing the antigenic proteins or peptides, in the manufacture of an agent for induction of antigen-specific T cells. The products of the invention have virucide and cytostatic activity and can be used for gene therapy or as inducers of antigen-specific T cells. The action of interferon-alpha (IFN-alpha) in a system for inducing specific cytotoxic real (CTL) by administering an antigenic peptide in an incomplete recell (CTL) by administering an antigenic peptide in an incomplete of recell (CTL) by administering an incomplete of received in the manufacture of a medicament for inducing antigen-specific T cells in an individual who has been administered with AP (or DNA encoding AP) or vice versa. The medicament is useful for the treatment or prophylaxis of a tumor or a viral infectious disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Use of antigenic proteins, peptides, interferon or their encoding DNA, in the manufacture of an agent for the induction of antigen-specific T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                   Tumor antigenic peptide; interferon; IFN; antigen-specific T cell; virucide; cytostatic; gene therapy; IFN-alpha; cytotoxic T cell; viral infectious disease; cyclophilin B; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .,
                                                                                                                                                                                                                   Human cyclophilin B derived tumor antigenic peptide SEQ ID 17.
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Pred. No. 1.7e+06;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAG68089 standard; peptide; 9 AA.
                                                                                                                  AAB46931 standard; peptide; 9 AA
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                                                                                                                                                                                    04-MAY-2001 (first entry)
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Best Local Similarity
DXMIQGGD
                              DYMIQGGD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-JUL-1999;
                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                        EP1074267-A1
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                                                                                                                                                    AAB46931;
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ID AAG6
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0; Gaps

Score 37; DB 4; Length 9; Pred. No. 1.7e+06; 0; Mismatches 1; Indels

h 92.5%; Similarity 87.5%; 7; Conservative

Query Match Best Local Similarity Matches 7; Conserv

Novel composition for treating a disease in an animal, comprises an immune effector cell and cell penetrating peptide associated with an

(BAYU) BAYLOR COLLEGE MEDICINE.

WPI; 2002-627577/67

Wang R;

antigen or antibody

15~FEB-2002; 2002WO-US005212. 15-FEB-2001; 2001US-0268687P.

WO200264057-A2.

22-AUG-2002

Homo sapiens.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes peptides recognising cancer cells with tumour specific cytotoxic T lymphocytes (CTL). The peptides recognising cancer cells with tumour specific CTLs are selected from: (1) peptides of sequences (AAG68066 to AAG68069); (2) peptides containing the above mentioned sequences; (3) peptides having 70 % or more of homogeneity with the above mentioned sequences; and (4) peptides with one or more deleted, substituted, added or inserted amino acid(s) of the above mentioned sequences, particularly those having recognising property due to HLA-A2402 binding CTL, especially having at least 5 amino acids, used for medicine, particularly anticancer agents, derived from antitumour antigenic peptides of 1ck, src family, SART-1, SART-3 or cyclophilin B genes. The antitumour peptides have cytostatic activities. The peptides are used for the treatment of cancer. The peptides cause activation of CTL in cancer patients. The present sequence represents a peptide from
                                                                                                                                                                                                                                                                                                                                                                                                                             New peptides for recognizing cancer cells with tumor specific cytotoxic Tlymphocytes and for treating cancer.
                                                                                                Antitumour; cancer; cancer cell recognition; antigenic; CTL; lck; src; tumour specific cytotoxic T lymphocyte; anticancer; SART-1; SART-3; cyclophilin B gene; HLA-A2402.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cell penetrating peptide; cancer; tumour; melanoma; thymoma; antigen; lymphoma; sarcoma; lung cancer; non-Hodgkin's lymphoma; leukaemia; Hodgkin's lymphoma; uterine cancer; cervical cancer; bladder cancer; kidney cancer; adenocarcinoma; breast cancer; prostate cancer; ovarian cancer; pancreatic cancer; epitope; vaccine; dendritic cell; tumour infiltrating lymphocyte; TIL; human leukocyte antigen; HLA;
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Pred. No. 1.7e+06;
0; Mismatches 1; Indels
                                                                   Antitumour peptide cyclophilin B 91-99.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 8; Page 2; 14pp; Japanese.
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87.5%;
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                                                                                                                                                                                                                                                                                                                     99JP-00374322
                                  (first entry)
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Best Local Similarity 87.5
Matches 7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 DXMIQGGD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DEMIQGGD
                                                                                                                                                                                                              JP2001245675-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                       (ITOY/) ITO Y.
                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                      28-DEC-1999;
                                  17-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-NOV-2002
                                                                                                                                                                                                                                                  11-SEP-2001
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AAG68089;
                                                                                                                                                                                                                                                                                                                                                                                                                                New
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 8
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The invention relates to a composition (I) comprising an immune effector cell and a cell penetrating peptide (CPP) associated with an antigen or antibody. Also included are (1) a vaccine comprising (I), CPP associated with an antigen, and a pharmaceutically acceptable carrier and (2) preparing a composition for a disease, by providing (I) and CPP associated with an antigen for disease, and introducing the antigeners of associated CPP to (I), where antigen enters into the cell. The antigens are, for example, tumour antigen derived epitopes recognised by tumour control of infiltrating lymphocytes (TIL) of HLA (human leukocyte antigen) class I or II. The composition is useful for enhancing immunity in an animal to a disease, by administering a mature dendritic cell comprising CPP associated with an antigen to disease, to the animal, such that following the administration, animal is protected from disease, where the animal comprises both CD4+ and CD8+ T cells. It is also useful for treating a disease (e.g. cancer, tumour, melanoma, thymoma, lymphoma, sarcoma, lung cancer, non-Hodgkin's lymphoma, leukaemia, Hodgkin's lymphoma, uterine cancer, non-Hodgkin's lymphoma, leukaemia, gancer, adenocarcinoma, haset cancer, and cancer, kidney cancer, adenocarcinoma, cancer, and cancer, adenocarcinoma, cancer, and cancer, adenocarcinoma, cancer, and cancer, adenocarcinoma, cancer, and cancer, adenocarcinoma, cancer, and cancer, adenocarcinoma, cancer, accertael cancer, and cancer, and cancer, and cancer, adenocarcinoma, cancer, adenocarcinoma, cancer, and cancer, and cancer, and cancer, and cancer, adenocarcinoma, and cancer, and cancer, and cancer, adenocarcinoma, cancer, and cancer, adenocarcinoma, and cancer, and canc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          breast cancer, prostate cancer, ovarian cancer and pancreatic cancer). The animal is further subjected to a cancer treatment including surgery, radiation, chemotherapy or gene therapy. The administration of (1), preferably dendritic cell is prior to, subsequent to or concurrent with, the cancer treatment. The present sequence is a tumour antigen derived epitope for inclusion in the composition of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          92.5%; Score 37; DB 5; Length 9; 87.5%; Pred. No. 1.7e+06; iive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human CypB HLA-A24 epitope, SEQ ID NO:24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 17; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR84374 standard; peptide; 9 AA.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 DXMIQGGD
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The invention relates to allergic reaction suppressors, comprising a peptide which originates from the same antigenic substance as the antigenic substance inducing the (peptide-originated) allergic reaction, and containing an epitope different from the epitope participating in the induction of the allergic reaction and yet dos not induce the allergic reaction. The desensitizers or allergic reaction suppressors are useful for treating and preventing type I allergic diseases and in cancer vaccines for preventing or treating cancer. The allergic reaction suppressors are also useful for suppressing an allergic reaction or reducing immunoglobulin E antibody production, and for desensitization to antigens. This sequence represents a peptide used in the invention and corresponds to amino acids 91-99 of the cyclophilin B protein with the phe at position 92 replaced by a Tyr.
                                               Desensitizers or allergic reaction suppressors containing peptides originating from the same antigenic substances as reaction-inducing substances, useful for preventing or treating type I allergic diseases
                                                                                                                                                                                                                                                                                                                                                                                                                   92.5%; Score 37; DB 7; Length 9; 87.5%; Pred. No. 1.7e+06; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cyclophilin B protein amino acids 91-99.
                                                                                                                            Claim 8; SEQ ID NO 9; 49pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 8; SEQ ID NO 2; 49pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC17702 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-AUG-2002; 2002WO-JP008641
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    7; Conservative
                                                                                                 and in cancer vaccines.
                  WPI; 2003-300831/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 DXMIQGGD 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-300831/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yamada A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 DYMIOGGD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2003020306-A1.
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                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC17702;
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                             The invention relates to a method for the detection of antigen specific T -cells in a blood sample involving the use of a plurality of antigenic peptides. The method comprises sampling of peripheral blood monocytes; stimulation of the collected peripheral blood monocytes with antigens without direct use of antigen presenting cells; and detection of T-cells specific to the antigen in the stimulated monocytes. The method is particularly used for the detection of cancer as it can be used in semi-quantitative determination of cancer specific T-cells. It can also be used for cancer vaccine therapy for patients with cervical or prostate cancer. The method can additionally be used to monitor of cellular in minimal the semi-cancer. The method can additionally be used to monitor of cellular in minimal than the semi-cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytostatic; antiallergic; vaccine; allergic reaction suppressor; antigen; allergy; epitope; cancer; immunoglobulin E antibody; desensitization; cyclophilin B.
                                                                                                                                                                                                                     A detection method of antigen specific T-cells, comprises the use of plural antigenic peptides, useful in semi-quantitative determination of cancer specific T-cell frequencies and for monitoring cellular immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunity and cancer immune therapy by detection of specific T-cell
frequencies. Sequences ABR84367-ABR84380 represent HLA-A24 (human
leukocyte antigen) peptides of human origin used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Pred. No. 1.7e+06;
); Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified cyclophilin B protein amino acids 91-99.
                                                                                                                                                                                                                                                                                      Example 8; Page 10; 18pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC17709 standard; peptide; 9 AA.
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87.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-AUG-2002; 2002WO-JP008641.
                                                                                             18-SEP-2001; 2001JP-00283413
                                                                                                                          13-NOV-2000; 2000JP-00345094
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Best Local Similarity
                                                                                                                                                                                         WPI; 2003-508315/48.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 DXMIQGGD
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                               JP2002365286-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 9 AA;
 Homo sapiens
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                                                              18-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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Gaps

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cytostatic; antiallergic; vaccine; allergic reaction suppressor; antigen; allergy; epitope; cancer; immunoglobulin E antibody; desensitization; cyclophilin B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to allergic reaction suppressors, comprising a
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Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.

Human cyclophilin B peptide fragment #37

(first entry)

11-APR-2000

AAY69957;

AAY69957 standard; peptide; 9 AA.

DEMIQGGD 8

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RESULT 13
                                                   AAY69957
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peptide which originates from the same antigenic substance as the antigenic substance inducing the (peptide-originated) allergic reaction, and containing an epitope different from the epitope participating in the induction of the allergic reaction and yet dos not induce the allergic reaction. The desensitizers or allergic reaction suppressors are useful for treating and preventing type I allergic diseases and in cancer vaccines for preventing or treating cancer. The allergic reaction suppressors are also useful for suppressing an allergic reaction or reducing immunoglobulin B antibody production, and for desensitization to antigens. This sequence represents a peptide used in the invention and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying a cell that differentiates into a metastatic cancer cell, useful for preventing metastatic cancer, comprises identifying a mutated fibronectin in the cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino
                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention comprises a method for identifying a cell that will differentiate into a metastatic cancer cell, the method involves identifying a mutated fibronectin in the cell. The method of the invention is useful for preventing metastatic cancer. The present acid sequence represents a Class I HLA-restricted widely expressed
                                                                                                                                                                                                          .;
                                                                                                                                corresponds to amino acids 91-99 of the cyclophilin B protein
                                                                                                                                                                                                                                                                                                                                                                                                                           metastatic cancer cell differentiation; mutated fibronectin; metastatic cancer; class I HLA-restricted; widely antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 37; DB 8; Length 9;
Pred. No. 1.7e+06;
0; Mismatches 1; Indels
                                                                                                                                                                                                         1; Indels
                                                                                                                                                                                Score 37; DB 7; Length 9; Pred. No. 1.7e+06; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                    Class I HLA-restricted widely expressed antigen #7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; SEQ ID NO 85; 137pp; English.
                                                                                                                                                                                                                                                                                                                          ADG89642 standard; peptide; 9 AA.
                                                                                                                                                                                                          ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BAYU ) BAYLOR COLLEGE MEDICINE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-MAY-2002; 2002US-0383530P.
                                                                                                                                                                                92.5%;
87.5%;
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                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                    Query Match
Best Local Similarity 8/...
7; Conservative
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                                                                                                                                                                                                                                   DXMIQGGD 8
                                                                                                                                                                                                                                                           DFMIQGGD 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 9 AA;
                                                                                                                                                         Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified
                                                                                                                                                                                                                                                                                                                                                                             11-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     useful
                                                                                                                                                                                                                                                                                                   RESULT 12
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Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.

(SUMU) SUMITOMO PHARM CO LTD (ITOH/) ITOH K.

WPI; 2000-116932/10.

Gomi S;

Itoh K,

99WO-JP003360. 98JP-00178449.

24-JUN-1999;

29-DEC-1999

25-JUN-1998;

Homo sapiens WO9967288-A1 Claim 4; Page 61; 64pp; Japanese.

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This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                             Rb-interacting zinc finger; RIZ; retinoblastoma; heart disease; cell proliferation; cell differentiation; tissue repair; transcription regulator; breast cancer; gene therapy; melanoma; neuroblastoma; leukaemia; Parkinson's disease; Huntingdon's disease; Alzheimer's disease; paralysis; motor neurone disorder; cel.
                                                                                                                                              Gaps
                                                                                                                                              0;
                                                                                                               Length 9;
                                                                                                               Score 27; DB 3; 1 Pred. No. 1.7e+06;
                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                           AAB12089 standard; peptide; 8 AA.
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                                                                                                                67.5%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unidentified adenovirus
                                                                                                                Query Match
Best Local Similarity
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                                                                                                                                                                              3 MIQGGD
                                                                                                                                                                                                                                                                                                                                                                                                    Ad7 cel peptide.
                                                                                                                                                                                                              MCOGGD
                                                                                  Sequence 9 AA;
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                                                                                                                                                                                                                                                                                                                                                                       17-JAN-2001
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                                                                                                                                                                                                                                                                                                                          AAB12089;
                                                                                                                                                Matches
                                                                                                                                                                                                                                                             RESULT 14
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Gaps

0

Query Match
Best Local Similarity 87.5%;
Matches 7; Conservative 0

DXMIQGGD

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Query Match 62.5%; Score 25; DB 3; Length 8;
Best Local Similarity 37.5%; Pred. No. 1.7e+06;
Matches 3; Conservative 4; Mismatches 1; Indels 0; Gaps 0;
Qy 1 DXMIQGGD 8
CDb 1 EDLLEGGD 8
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cytostatic; tumor; gene therapy; retinoblastoma protein-interacting zinc protein; RIZ; neuroblastoma; melanoma; cardiac cell growth inducer; neuronal cell growth inducer;

adenovirus; ElA; cel. unidentified adenovirus.

JS6468985-B1

Adenovirus 7 E1A gene cel domain.

(first entry)

29-JAN-2004

ADE84662;

ADE84662 standard; peptide; 8 AA.

RESULT 15 ADE84662

Job time : 86 secs

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0
                                                                                                                                                                                                                                                   Reducing the growth of tumor cell having mammalian retinoblastoma protein -interacting zinc finger proteins in a subject, comprises administering nucleic acid molecule encoding RIZ at or adjacent to the site of tumor.
                                                                                                                                                                                                                                                                                                                                                     The invention describes a method of reducing growth of a tumour cell having a mutant mammalian retinoblastoma (Rb) protein-interacting zinc finger (RIZ) protein in a subject The method comprises administering a nucleic acid molecule (I) encoding RIZ at or adjacent to tumour site. (I) is administered at the site of the tumour. The method is useful for reducing the growth of a tumour cell e.g. neuroblastoma or melanoma cell having mutant RIZ protein. The method is useful for inducing growth of a cardiac cell or a neuronal cell in a subject, and for effecting normal growth control to a tumour cell or causing differentiation of tumour cells. This is the amino acid sequence of an adenovirus crl domain used in a comparison with the cel domain of RIZ protein.
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Pred. No. 1.7e+06;
4; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                        Example 3; SEQ ID NO 75; 49pp; English.
                                                                     94US-00292683.
95US-00399411.
95US-00516859.
                                  17-MAR-2000; 2000US-00528706.
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                                                                                                                                           (BURN-) BURNHAM INST
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Best Local Similarity
Matches 3; Conserv
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                                                                    18-AUG-1994;
06-MAR-1995;
18-AUG-1995;
22-OCT-2002
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us-09-720-469a-38.rai

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3; Conservative
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amino acid
GY: linear
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CITY: San Diego
STATE: California
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Best Local Similarity
Matches 3; Conserv
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Sequence 84, Appl
Sequence 84, Appl
Patent No. 5196404
Patent No. 5433940
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                                                             October 21, 2004, 07:13:10 ; Search time 21.5 Seconds (without alignments) 27.761 Million cell updates/sec
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Sequence 62, A
Sequence 1, Ap
Sequence 1, Ap
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Sequence 5,
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Sequence 75
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Sequence
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2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
        GenCore version 5.1.6
(c) 1993 - 2004 Compugen Ltd.
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US-08-399-411-75
US-08-516-859A-75
US-09-586-472-75
US-09-528-706-75
US-08-457-172-1
US-08-457-172-1
US-08-953-327-1
US-08-215-805A-7
US-08-215-805A-52
US-08-215-805A-58
US-08-215-805A-58
US-08-215-805A-19
US-07-620-426B-19
US-07-620-426B-19
US-07-620-426B-19
US-07-620-426B-19
US-07-620-426B-19
US-08-215-805A-48
US-08-215-805A-48
US-08-215-805A-48
US-08-215-805A-48
US-08-215-805A-44
US-08-215-805A-44
US-08-215-805A-44
US-08-215-805A-44
US-08-215-805A-44
US-08-215-805A-44
US-08-215-805A-44
US-08-215-805A-44
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US-08-633-779-3
US-08-633-779-5
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Maximum Match 100%
Listing first 45 summaries
                                              OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Match Length DB
                                                                                                                           1 DXMIQGGDX 9
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Maximum DB seq length: 12
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Sequence 227, App
Sequence 101, Appl
Sequence 13, Appl
Sequence 13, Appl
Patent No. RE34,60
Sequence 39, Appl
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Sequence 39,
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TITLE OF INVENTION: Retinoblastoma Protein - Interacting
TITLE OF INVENTION: Zinc Finger Proteins
NUMBER OF SEQUENCES: 93
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                62.5%; Score 25; DB 2; Length 8; 37.5%; Pred. No. 3.8e+05; ive 4; Mismatches 1; Indels
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ZIP: 92122
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Batentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/459,568
FILING DATE: 02-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/399,411
FILING DATE: 06-MAR-1995
ATTORNEY/AGENT INFORMATION:
FILING DATE: 06-MAR-1995
ATTORNEY/AGENT INFORMATION:
REFERENCE/DOCKET NUMBER: P-LJ 1264
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 75:
SEQUENCE CHARACTERISTICS:
LENGTH CHARACTERISTICS
LENGT
US-09-846-350A-1

US-09-569-037-1

US-08-329-820-227

US-08-329-820-101

US-08-72-256-13

US-09-111-696B-13

RE34606-25

US-07-940-861-39

US-08-459-512-39

US-08-459-512-39

US-08-244-496-70

US-08-244-496-70

US-08-459-657-39

US-08-459-657-39
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Patent No. 5811304
GENERAL INFORMATION:
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US-08-516-859A-75
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                                   Sequence 75, Application US/08399411

Sequence 75, Application US/08399411

Patent No. 5831008

GENERAL INFORMATION:

APPLICANT: Huang, Shi

TITLE OF INVENTION: Retinoblastoma Proteins

TITLE OF INVENTION: Zinc Finger Proteins

NUMBER OF SEQUENCES: 93

CORRESPONDENCE ADDRESS:

ADDRESSEE: Campbell and Flores

STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 75. Application US/08516859A
Sequence 75. Application US/08516859A
Patent No. 6069231
GENERAL INFORMATION:
APPLICANT: Huang, Shi
TITLE OF INVENTION: Retinoblastoma Protein - Interacting
TITLE OF INVENTION: Zinc Finger Proteins
NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                62.5%; Score 25; DB 2; Length 8; 37.5%; Pred. No. 3.8e+05;
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/399,411
FILING DATE: 06-MAR-1995
CLASSIFICATION: 530
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/516,859A
FILING DATE: 18-AUG-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 1264
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 75:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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Best Local Similarity 37.5
----hes 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: San Diego
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                          CITY: San Diego
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
TOPOLOGY: linear
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EDLLEGGD 8
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US-08-516-859A-75
RESULT 2
US-08-399-411-75
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Sequence 75, Application US/09586472
Patent No. 6323335
GENERAL INFORMATION:
APPLICANT: Huang, Shi
TITLE OF INVENTION: Retinoblastoma Protein - Interacting
Zinc Finger Proteins
NUMBER OF SEQUENCES: 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 25; DB 3; Length 8; Pred. No. 3.8e+05; 1; Indels 4; Mismatches 1; Indels
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
CITY: San Diego
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/586,472
FILING DATE: 01-Jun-2000
CLASSIFICATION: <Unknown>
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APPLICATION NUMBER: US 09/528,706
FILING DATE: 17-MAR-2000
APPLICATION NUMBER: US 08/516,859
FILING DATE: 18-AUG-1995
APPLICATION NUMBER: US 08/399,411
FILING DATE: 06-MAR-1995
APPLICATION NUMBER: US 08/292,683
FILING DATE: 18-AUG-1994
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PRIOR APPLICATION: 314

PRIOR APPLICATION DATA:

#PRIOR APPLICATION NUMBER: US 08/399,411

FILING DATE: 06-MAR-1995

PRIOR APPLICATION DATA:

#PRIOR APPLICATION NUMBER: US 08/292,683

#PRIOR APPLICATION NUMBER: B-LJ 1776

#PRIOR APPLICATION NUMBER: P-LJ 1776

#PRIOR APP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
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Similarity 37.5%;
3; Conservative '
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SEQUENCE CHARACTERISTICS
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COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8 amino acids
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1 EDLLEGGD 8
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Best Local Similarity
Matches 3; Conserv
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3 MIQGGD 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 7
US-08-487-860-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 δ
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Sequence 75, Application US/09528706
GENERAL INFORMATION:
APPLICANT: Huang, Shi
TITLE OF INVENTION: Retinoblastoma Protein - Interacting
TITLE OF INVENTION: Zinc Finger Proteins
NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                              Score 25; DB 3; Length 8; Pred. No. 3.8e+05; 4; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  62.5%; Score 25; DB 4; Length 8; 37.5%; Pred. No. 3.8e+05; tive 4; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 9212.

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/528,706
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:

PRIOR APPLICATION NUMBER: US 08/292,683

FILING DATE: 18-AUG-1994

ATTORNEY/AGENT INFORMATION:

NAME: Campbell, Cathryn A.

REGISTRATION NUMBER: 31,815

REFERENCE/DOCKET NUMBER: P-LJ 1776

TELECOMMUNICATION INFORMATION:

TELEPHONE: (619) 535-9001

TELEFAX: (619) 535-8949

INFORMATION FOR SEQ ID NO: 75:

SEQUENCE CHARACTERISTICS:
TYPE: amino acid
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 75:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/516,859
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 1, Application US/08457172 ; Patent No. 5622837
                                                                                               62.5%;
                                                                                         Query Match 62.53
Best Local Similarity 37.53
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 8 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: San Diego
STATE: California
COUNTRY: USA
                                                                                                                                                                         1 DXMIQGGD 8
                                                                                                                                                                                                  : ::: | | |
1 EDLLEGGD 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 DXMIQGGD 8
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EDLLEGGD 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                        US-09-586-472-75
                                                                                                                                                                                                                                                                                           US-09-528-706-75
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US-08-457-172-1
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GENERAL INFORMATION:
APPLICANT: Hans Scheefers, Ursula Scheefers-Borchel and
APPLICANT: Hans Scheefers, Ursula Scheefers-Borchel and
APPLICANT: Andreas Sziegoleit
TITLE OF INVENTION: PANCREAS ELASTASE 1-SPECIFIC ANTIBODY, A PROCESS FOR
TITLE OF INVENTION: OBTAINING IT, AND A TEST KIT CONTAINING SUCH ANTIBODY
NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 62, Application US/08487860
; Patent No. 5792456
; GENERAL INFORMATION:
    APPLICANT: Yelton, Dale
    APPLICANT: Huse, William
    APPLICANT: Rosok, Mae J.
    TITLE OF INVENTION: No. 5792456el Mutant BR96 Antibodies and
    TITLE OF INVENTION: Punctional Equivalents Reactive With Human Carcinomas
    NUMBER OF SEQUENCES: 62
    CORRESPONDENCE ADDRESS:
    ADDRESSEE: Merchant & Gould
    STREET: 11150 Santa Monica Bldv., Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Pred. No. 3.8e+05;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 42883/JPW/AJM TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 977-9550 TELEFAX: (212) 977-9809 TELEX: 422523 COOP UI INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                              ZIP: 10112
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/457,172
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/220,395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 07/969,173
FILING DATE: 29-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
                                                                                                                                                             ADDRESSEE: Cooper & Dunham STREET: 30 Rockefeller Plaza CITY: New York STATE: New York COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              62.5%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ouery Match
Best Local Similarity 66...
4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CELL TYPE: pancreatic US-08-457-172-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  unknown
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CA
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CLESSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/953,327
FILING DATE:
APPLICATION NUMBER: 60/028,701
FILING DATE: 18-OCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: Saliwanchik, David R.
REGISTRATION NUMBER: 31,794
REFERENCE/DOCKET NUMBER: UF-171
TELECOMMUNICATION INFORMATION:
TELEPHONE: (352) 375-8100
TELEFAX: (352) 375-8100
TELEFAX: (352) 375-5800
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                    Query Match
Best Local Similarity 62.5
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ) TOPOLOGY: linear
) MOLECULE TYPE: peptide
US-09-472-579-1
                                                             TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-953-327-1
                                            single
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                                                                                                                                                                                                                                        1 DXMIQGGD 8
                      TYPE: amino acid
STRANDEDNESS: sin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 10
US-08-215-805A-7
LENGTH:
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Sequence 1. Application US/08953327
Sequence 1. Application US/08953327
GENERAL INFORMATION:
APPLICANT: Staud, Roland
TITLE OF INVENTION: AND TREATMENT OF WEGENER'S GRANULOMATOSIS
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: Saliwanchik & Saliwanchik
STREET: 2421 N.W. 41st Street, Suite A-1
CITY: Gainesville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .
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     COUNTRY: USA
ZIP: 90025-3395
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/487,860
FILING DATE: 07-JUN-1995
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Adriano, Sarah B.
REFERENCE/DOCKET NUMBER: 34,470
REFERENCE/DOCKET NUMBER: 34,470
TELEPHONE: 310-445-1140
TELEFAX: 310-445-9031
INFORMATION FOR SEQ ID NO: 62:
SEQUENCE CHARACTERISTICS:
'.FNATH: 8 amino acids
'.FNATH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: FL.
COUNTRY: US
ZIP: 32606-6669
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/953,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/028,701
FILING DATE: 18-OCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: Saliwanchik, David R.
REFERENCE/DOCKET NUMBER: 31,794
REFERENCE/DOCKET NUMBER: UF-171
TELECOMMUNICATION INFORMATION:
TELEPHONE: (352) 375-8100
TELEPAX: (352) 375-8100
TELETAX: (352) 375-8100
TELEPAX: (352) 375-8100
TELEPAX: (352) 375-8100
TELEPAX: (352) 375-8100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , MOLECULE TYPE: peptide US-08-487-860-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 QGGD 8
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   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 8
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                                                     Gaps
                                                                                                                                                                                                                                                  US-09-472-579-1
; Sequence 1, Application US/09472579
; Patent No. 6277955
; GENERAL INFORMATION:
; TITLE OF INVENTION: MATERIALS AND METHOD FOR THE DETECTION
; TITLE OF INVENTION: AND TREATMENT OF WEGENER'S GRANULOMATOSIS
; NUMBER OF SEQUENCES: 3
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Saliwanchik & Saliwanchik
; STREET: 2421 N.W. 41st Street, Suite A-1
; CITY: Gainesville
                                                     ..
O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
57.5%; Score 23; DB 3; Length 11; 62.5%; Pred. No. 1.9e+02; ive 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     57.5%; Score 23; DB 3; Length 11; 62.5%; Pred. No. 1.9e+02; Live 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTRY: US
ZIP: 32606-6669
ZIP: 32606-6669
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/472,579
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Conservative
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MOLECULE TYPE: peptide
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ::[||
1 LEGGD 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
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Sequence 7, Application US/08215805A
Fatent No. 5559008
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES: 84
CORRESPONDENCE ADDRESS:
ADDRESSE: Nixon, Hargrave, Devans & Doyle
STREET: Clinton Square, P.O. Box 1051
CITY: Rochester
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USOUR 11.

USOUR 211.

USOUR 211.

USOUR 212.

Sequence 52, Application US/08215805A

Patent No. 5559008

GENERAL INFORMATION:

TITLE OF INVENTION: LEUKOTOXIN GENE FROM PASTEURELLA

TITLE OF INVENTION: SUIS

NUMBER OF SEQUENCES: 84

CORRESPONDENCE ADDRESS:

ADDRESSEE: Nixon, Hargrave, Devans & Doyle

STREET: Clinton Square, P.O. Box 1051

CITY: Rochester

STATE: New York

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      55.0%; Score 22; DB 1; Length 9; 80.0%; Pred. No. 3.8e+05;
                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.30
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/215,805A
FILING DATE: 22-MAR-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
RAME: Timian, Susan J.
REGISTRATION NUMBER: 34,103
REFERENCE/DOCKET NUMBER: 19603/61 (D-1329A)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (716) 263-1636
TELEFAX: (716) 263-1600
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LIBRARY: P. suis DNA in Bacteriophage lambda-dash CLONE: (Lambda)yfc33-37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 14603
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Pasteurella suis
STRAIN: 5943
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: peptide
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 IQGGD 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 IDGGD 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-215-805A-7
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Gaps
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Sequence 58, Application US/08215805A

Patent No. 5559008

GENERAL INFORMATION:
TITLE OF INVENTION: LEUKOTOXIN GENE FROM PASTEURELLA

TITLE OF INVENTION: SUIS

NUMBER OF SEQUENCES: 84

CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon, Hargrave, Devans & Doyle
STREET: Clinton Square, P.O. Box 1051

CITY: Rochester
STATE: New York

COUNTRY: USA

ZIP: 14603

ZIP: 14603

ZIP: 14603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/215,805A
FILING DATE: 22-MAR-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Timian, Susan J.
REGISTRATION NUMBER: 34,103
REFERENCE/DOCKET NUMBER: 19603/61 (D-1329A)
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF SEQ ID NO: 58:
SEQUENCE CHARACTERISTICS:
T.FN/7TH: 9 amino acids
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/215,805A
FILING DATE: 22-MAR-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Timian, Susan J.
REGISTRATION NUMBER: 19603/61 (D-1329A)
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (716) 263-1600
INFORMATION FOR SEQ ID NO: 52:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
TYPE: amino acids
STRANDEDNESS:
STR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide
CRIGINAL SOURCE:
CREANISM: Actinobacillus pleuropneumoniae
US-08-215-805A-52
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Query Match
Best Local Similarity
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US-07-662-007B-19
                                                                                                                                                                                                                                                           US-07-620-426B-19
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GENERAL INFORMATION:

APPLICANT: Choi, Eui-Sung
APPLICANT: Rhee, Sang-Ki
APPLICANT: Lee, Eun-Hae
TITLE OF INVENTION: and Methods of Use Thereof
FILE REFERENCE: 1533.0870000
CURRENT FILING DATE: 1999-04-22
NUMBER OF SEQ ID NOS: 87

SOFTWARE:

APPLICANT: APPLICATION NUMBER: US/09/296,284A

CURRENT FILING DATE: 1999-04-22

SOFTWARE:

SOFTWARE:

AUGUST PATENTIN Ver. 2.0
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Pred. No. 4.2e+02;
1; Mismatches 2; Indels
                                                            Length 9;
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ZIP: 95616
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 720 KB storage COMPUTER: Apple Macintosh OPERATING SYSTEM: Macintosh 6.0.7
SOFTWARE: MicrosoftWord 4.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/620,426B
FILING DATE: 19901130
                                                           Query Match 55.0%; Score 22; DB 1; I Best Local Similarity 80.0%; Pred. No. 3.8e+05; Matches 4; Conservative 0; Mismatches 1;
; ORGANISM: Actinobacillus pleuropneumoniae US-08-215-805A-58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19, Application US/07620426B; Patent No. 5298421; GENERAL INFORMATION:
APPLICANT: Davies, Huw Maelor
APPLICANT: Pollard, Michael Roman
APPLICANT: Voekler, Toni Alois
TITLE OF INVENTION: Plant Thioesterases
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/514,030
FILLING DATE: 26-APRIL-1990
ATTORNEY/AGENT INFORMATION:
NAME: Elizabeth Lassen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Gluconobacter suboxydans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          52.5%;
57.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Calgene, Inc.
STREET: 1920 Fifth Street
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Best Local Similarity 57.1
Matches 4; Conservative
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STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 DXMIQGG 7
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LENGTH: 10
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COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.50 inch, 720 KB storage COMPUTER: Apple Macintosh
OPERATING SYSTEM: Macintosh 6.0.7
SOFTWARE: MicrosoftWord 4.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/662,007B
FILING DATE: 19910408
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/514,030
FILING DATE: 26-APR-1990
ATTORNEY/AGENT INFORMATION:
NAME: Elizabeth Lassen
REGISTRATION NUMBER: 31,845
REFERENCE/DOCKET NUMBER: 31,845
REFER
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APPLICANT: Pollard, Michael Roman
APPLICANT: Voekler, Toni Alois
APPLICANT: Thompson, Gregory A.
TITLE OF INVENTION: Plant Thioesterases
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
REGISTRATION NUMBER: 31,845
NAME: Donna E. Scherer
REGISTRATION NUMBER: 34,719
REFERENCE/DOCKET NUMBER: CGNE 70-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 916-753-6313
TELEFAX: 916-753-1510
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19, Application US/07662007B; Patent No. 5344771
GENERAL INFORMATION:
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1920 Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                , MOLECULE TYPE: peptide US-07-662-007B-19
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STATE: California
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0; Gaps

Matches 4; Conservative 0; Mismatches 0; Indels

4 IQGG 7 |||| 3 IQGG 6

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Search completed: October 21, 2004, 07:29:29 Job time : 22.5 secs

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Sequence 85, Appli
Sequence 2, Appli
Sequence 9, Appli
Sequence 9, Appli
Sequence 14, Appl
Sequence 161, Appl
Sequence 161, Appl
Sequence 124, Appl
Sequence 124, Appl
Sequence 30, Appl
Sequence 31, Appl
Sequence 35, Appl
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47.187 Million cell updates/sec
                                                            October 21, 2004, 07:15:25 ; Search time 61.75 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Description
                                                                                                                                                                                                                                                                                                    GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-10-788-016-2

US-10-788-016-9

US-10-128-520-96

US-10-210-148-9

US-10-210-148-14

US-10-316-35-32

US-10-356-257-161

US-10-620-052A-50

US-09-765-086-84
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US-10-014-326-21
US-10-264-374-84
                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                                             1364641 seqs, 323758627 residues
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                                                                                                                                                                                                                                                                                              Published Applications AA:*
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Maximum Match 100%
Listing first 45 summaries
                                       OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Match Length DB
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	equence 28 equence 98 equence 98 equence 44 equence 57 equence 11 equence 27 equence 27
	10-723-933-8 10-723-933-8 10-462-452-4 10-601-953-5 10-322-266-4 10-723-933-2 10-723-933-8
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ALIGNMENTS

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                                                                              APPLICANT: Wang, Rong-fu
TITLE OF INVENTION: Mutant Fibronectin and Tumor Metastasis
FILE REFERENCE: HO-P02484US1
CURRENT APPLICATION NUMBER: US/10/447,161
CURRENT FILING DATE: 2003-05-28
PRIOR APPLICATION NUMBER: 60/383,530
PRIOR FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 148
SOFTWARE: Patentin version 3.1
SEQ ID NO 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                             92.5%; Score 37; DB 15; Length 9; 87.5%; Pred. No. 1.2e+06; Live 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Synthetic Peptide
                 Sequence 85, Application US/10447161 Publication No. US20040023314A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/10788016
Publication No. US20040141992A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 87.5
Matches 7; Conservative
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US-10-447-161-85
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US-10-447-161-85
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US-10-788-016-2
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KESULT 6
US-10-148-14
Sequence 14, Application US/10210148
Sequence 14, Application US/10210148
Sequence 14, Application No. US20030171280A1
GENERAL INFORMATION:
APPLICANT: Soderstrom, Karl Petter
TITLE OF INVENTION: Compositions And Methods For Mcdulation Of Immune Response
FILE REFERENCE: TROM0002
CURRENT APPLICATION NUMBER: US/10/210,148
CURRENT FILING DATE: 2002-07-31
PRIOR FILING DATE: 2002-07-31
PRIOR PILING DATE: 2002-07-31
NUMBER OF SEQ ID NOS: 117
SOFTWARE: Patentin Version 3.2
SEQ ID NO 14
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                US-10-210-148-9

Sequence 9, Application US/10210148

Sequence 9, Application US/10210148

Publication No. US20030171280A1

GENERAL INFORMATION:

TITLE OF INVENTION: Compositions And Methods For Mcdulation Of Immune Response

TITLE OF INVENTION: Compositions And Methods For Mcdulation Of Immune Response

CURRENT APPLICATION NUMBER: US/10/210,148

CURRENT FILING DATE: 2002-07-31

PRIOR APPLICATION NUMBER: PCT/US02/24311

PRIOR FILING DATE: 2002-07-31

NUMBER OF SEQ ID NOS: 117

SOFTWARE: PatentIn Version 3.2
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Pred. No. 1.2e+06;
1; Mismatches 1; Indels
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Pred. No. 9.7e+02;
3; Mismatches 2; Indels
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66.7%; Pred. No. 1.2e+06;
tive 1; Mismatches 1;
PRIOR APPLICATION NUMBER: US 60/285,683
PRIOR FILING DATE: 2001-04-24
NUMBER OF SEQ ID NOS: 381
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 96
LENGTH: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       52.5%;
66.7%;
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Best Local Similarity 66.7
Matches 4; Conservative
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US-10-210-148-14
                                                                                                                                  TYPE: PRT
CRGANISM: Homo sapiens
US-10-128-520-96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ) ORGANISM: Homo sapiens
US-10-210-148-9
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                                                                                                                                                                                                                                                                                                     1 DXMIQGGD 8
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Best Local Similarity
Matches 3; Conserv.
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Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Peptide consisting of 9 amino acid residues from the 91st residue; OTHER INFORMATION: to the 99th residue of cyclophillin B
US-10-788-016-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ) OTHER INFORMATION: Designed peptide based on the peptide consisting of 9 amino acid; OTHER INFORMATION: residues from the 91st residue to the 99th residue of; OTHER INFORMATION: cyclophillin B
US-10-788-016-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Indels
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Pred. No. 1.2e+06;
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87.5%; Pred. No. 1.-
.-- 0; Mismatches
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US-10-788-016-9
; Sequence 9, Application US/10788016
; Publication No. US20040141992A1
; GENERAL INFORMATION:
; APPLICANT: ITOH, Kyogo
; TITLE OF INVENTION: Desensitizers
; FILE REFERENCE: 3190-049
; CURRENT FILING DATE: 2004-02-26
; PRIOR FILING DATE: 2004-02-26
; PRIOR FILING DATE: 2001-08-29
; RIOR FILING DATE: 2001-08-29
; NUMBER OF SEQ ID NOS: 9
; SOFTWARE: PATENTIN VERSION 3.2
; LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 96, Application US/10128520; Publication No. US20040105862A1; GENERAL INFORMATION:
; GENERAL INFORMATION:
; TITLE OF INVENTION: Human TIMP-1 Antibodies; FILE REFERENCE: 02973.00073; CURRENT APPLICATION NUMBER: US/10/128,520; CURRENT FILING DATE: 2002-04-24
                                   FILE REFERENCE: 3190-049;
CURRENT APPLICATION NUMBER: US/10/788,016
CURRENT FILING DATE: 2004-02-26;
PRIOR APPLICATION NUMBER: PCT/JP02/08641;
PRIOR FILING DATE: 2002-08-28;
PRIOR FILING DATE: 2001-08-29;
PRIOR FILING DATE: 2001-08-29;
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.2;
                     TITLE OF INVENTION: Desensitizers
                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 87.5
Matches 7; Conservative
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                                                                                                                                                                                                                                                                    TYPE: PRT ORGANISM: Artificial
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US-10-128-520-96
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Query Match
Best Local Similarity
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LIEGG 5
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RESULT 9
US-10-076-047A-124
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APPLICANT: Schaeffer, Michael
APPLICANT: Schneiderbauer, Michael
APPLICANT: Weidler, Sascha
APPLICANT: Weidler, Sascha
APPLICANT: Hallek, Michael
APPLICANT: GSF Forschungszentrum fuer Umwelt und Gesundheit GmbH
TITLE OF INVENTION: Inhibition of the Proliferation of Cells of the
TITLE OF INVENTION: Multiple Myeloma
FILE REFERENCE: 080345-000100US
CURRENT APPLICATION NUMBER: US/10/194,985
CURRENT FILING DATE: 2002-07-12
PRIOR APPLICATION NUMBER: US 09/625,225
PRIOR FILING DATE: 2000-07-24
NUMBER OF SEQ ID NOS: 32
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 32
LENGTH: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15, Application US/10356257
; Publication No. US20040087767A1
; GENERAL INFORMATION:
; APPLICANT: LAZARUS, ROBERT A.
; APPLICANT: MAUN, HENRY R.
; TITLE OF INVENTION: FVIIA Antagonists
; FILE REFERENCE: P1950R1
; CURRENT APPLICATION NUMBER: US/10/356,257
; CURRENT FILING DATE: 2003-01-30
; PRIOR FILING DATE: 2002-02-06
; NUMBER OF SEQ ID NOS: 355
; SEQ ID NO 161
; LENGTH: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: sequence is synthesized
                                                                                                 Sequence 32, Application US/10194985
Publication No. US20030105287A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Artificial sequence
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Best Local Similarity 60.0
Matches 3; Conservative
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Sequence 124, Application US/10076047A; Sequence 124, Application US/10076047A; Publication No. US20030152935Al; Bublication No. US20030152935Al; GENERAL INFORMATION:

APPLICANT: Herath, Herath Mudiyanselage Athula Chandrasiri; TITLE OF INVENTION: Proteins, Genes and Their Use for TITLE OF INVENTION: Diagnosis and Treatment of Breast Cancer; FILE REFERENCE: 2543-1-026; CURRENT APPLICATION NUMBER: US/10/076,047A; CURRENT APPLICATION NUMBER: US 9919258.5; PRIOR APPLICATION NUMBER: GB 9919258.5; PRIOR FILING DATE: 2000-03-30; PRIOR FILING DATE: 2000-03-30; PRIOR FILING DATE: 2000-08-14; NUMBER OF SEQ ID NOS: 351; SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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; OTHER INFORMATION: Description of Artificial Sequence:consensus
; OTHER INFORMATION: peptide
US-10-620-052A-50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 50, Application US/10620052A; Sequence 50, Application US/10620052A; Publication No. US20040126784A1; Publication No. US20040126784A1; GENERAL INFORMATION:

APPLICANT: Hitoshi, Yasumichi
APPLICANT: Jenkins, Yonchu
APPLICANT: Markovtsov, Vadim
APPLICANT: Rigel Pharmaceuticals, Inc.
TITLE OF INVENTION: Modulators of Cellular Proliferation; FILE REFERENCE: 021044-004010US; CURRENT APPLICATION NUMBER: US/10/620,052A; CURRENT FILING DATE: 2003-07-14; PRIOR FILING DATE: 2002-07-12

NUMBER OF SEQ ID NOS: 78; SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              50.0%; Score 20; DB 14; Length 5; 60.0%; Pred. No. 1.2e+06;
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US-09-765-086-84
; Sequence 84, Application US/09765086
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                                                                                 APPLICANT: Waddening)
APPLICANT: Bredesen, Dale E.
APPLICANT: Ellerby, H. Michael
TITLE OF INVENTION: Chimeric Prostate-Homing Peptides With
TITLE OF INVENTION: Pro-Apoptotic Activity
TITLE OF INVENTION: P-LJ 3844
CURRENT APPLICATION NUMBER: US/09/765,086
CURRENT FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: US 09/489,582
PRIOR FILING DATE: 2000-01-21
NUMBER OF SEQ ID NOS: 235
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 35, Application US/09912787

Sequence 35, Application US/09912787

Patent No. US20020119149A1

GENERAL INFORMATION:

APPLICANT: JAKOBSEN, Bent Karsten

TITLE OF INVENTION: Multivalent T Cell Receptor Complexes

FILE REFERENCE: 102286.410DIV

CURRENT APPLICATION NUMBER: US/09/912,787

CURRENT FILING DATE: 2001-07-25

PRIOR FILING DATE: 1999-06-17

PRIOR APPLICATION NUMBER: PCT/GB99 01583

PRIOR FILING DATE: 1999-05-19

PRIOR FILING DATE: 1998-05-19

PRIOR FILING DATE: 1998-05-19

PRIOR FILING DATE: 1998-05-19

PRIOR FILING DATE: 1998-09-29

NUMBER OF SEQ ID NOS: 85

SOFTWARE: PatentIn Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       / Match 50.0%; Score 20; DB 9; I Local Similarity 50.0%; Pred. No. 1.2e+06; nes 3; Conservative 2; Mismatches 1;
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60.0%; Pred. No. 1.2e+06;
iive 1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: synthetic peptide
               GENERAL INFORMATION:
APPLICANT: Ruoslahti, Erkki
APPLICANT: Pasqualini, Renata
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ORGANISM: Artificial Sequence
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US20010046498A1
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Best Local Similarity
Matches 3; Conserv
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MVRDGD 8
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                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 84
LENGTH: 9
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 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
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MVDGG 5

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GS-Quence C. 10,14,236-21

GS-Quence C. 10,14,236-21

GS-Quence C. 10,1001(cont.com.18)/1001436

DSDD[CSRITION NO. UG207020142369A1

REPRICKAT: WILLOW UG207020142369A1

REPRICKAT: WILLOW C. GSO. GSOGGE P. G
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US-10-375-992-84
; Sequence 84, Application US/10375992
; Publication No. US20030152578A1
; GENERAL INFORMATION:
; APPLICANT: Ruoslahti, Erkki
; Pasqualini, Renata
; TITLE OF INVENTION: Tumor Homing Molecules, Conjugates
; TITLE OF INVENTION: Tumor Homing Molecules of Using Same
                                                     Gaps
                                                   .;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: United States

ZIP: 92122

COMPUTER EADABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/375,992

FILING DATE: 27-Feb-2003

CLASSIFICATION NUMBER: US/08/926,914

FILING DATE: 10-SEP-1997

ATTORNEY/AGENT INFORMATION:

NAME: Campbell, Cathryn A.

REGISTRATION NUMBER: 31,815

REFERENCE/DOCKET NUMBER: P-LJ 2725

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELEPHONE: (619) 535-9901

TELEFAX: (619) 535-9949

INFORMATION FOR SEQ ID NO: 84:

SEQUENCE CHARACTERISTICS:

LENGTH: 9 maino acids
Query Match 50.0%; Score 20; DB 14; Length 9; Best Local Similarity 50.0%; Pred. No. 1.2e+06; Matches 3; Conservative 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 50.0%; Score 20; DB 14; Length 9; Best Local Similarity 50.0%; Pred. No. 1.2e+06; Matches 3; Conservative 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 199
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores
STREET: 4370 La Jolla Village Drive, Suite 700
CITY: San Diego
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
MOLECULE TYPE: peptide
;
SEQUENCE DESCRIPTION: SEQ ID NO: 84:
US-10-375-992-84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
TOPOLOGY: both
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3 MVRDGD 8
                                                                                                 \delta
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Gaps

0;

Search completed: October 21, 2004, 07:33:42 Job time: 62.75 secs

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

October 21, 2004, 07:12:54; Search time 15.75 Seconds (without alignments) 54.981 Million cell updates/sec

US-09-720-469A-38 40 Title: Perfect score:

1 DXMIQGGDX 9 Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

283416 segs, 96216763 residues Searched:

1582 Total number of hits satisfying chosen parameters:

seq length: 0 seq length: 12

Minimum DB Maximum DB

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database

PIR 79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

kinesin light chai Ig H2 chain - Paci T-cell receptor be delta sleep-induci T-cell receptor be T-cell receptor ga Trimeresurus serin spinal cord peptid c-rel protein - ch triacylglycerol li Ig heavy chain CRD serum heterodimer, thymic factor - pi thymocyte growth p T cell receptor Vhypothetical prote T-cell receptor be hemoglobin, extrac hypothetical prote retinal oxidase -protein kinase C i glycoprotein hormo serine/threonine-s bradykinin-potenti orf 4 rara 5'-regi angiotensin-conver ubiquitin - rat globulin IV alpha Ig heavy chain DJ Description SUMMARIES S43170 B60769 PT0542 QDRB PN0044 PT0218 I41946 A61575 A23751 S57575 D37196 S19015 PH1458 A61309 PH1343 S60354 YFPG A60957 Query Match Length DB Score Result

-		7. Et
		ation of
	transforming prote T-cell receptor be protein QA300025 - calliMIRFamide 1 - pollen major aller cocoonase (BC 3.4. myosin light chain seed protein ws-5 T-cell receptor be T-cell receptor be T-cell receptor be NADH2 dehydrogenas neural cell adhesi quinoline 2-oxidor 3',5'-cyclic-GMP p neuropeptide Grb-A neuropeptide Grb-A	<pre>#text_change 09-Jul-2004 D. rization and chromosomal ID:g468786; PIDN:CAA49349</pre>
	A38590 S26544 PA0030 E44787 A53252 B61491 I50385 E61491 PT0667 PT0586 C39690 C39690 A53797 A53797 A57444	RESULT 1 843170 kinesin light chain - human (fragment) C;Species: Homo sapiens (man) C;Date: 13-Jan-1995 #sequence_revision 18-Aug-2000 C;Accession: 843170 R;Chernajovsky, Y.; Brown, A.; Jones, T.A.; Sheer, submitted to the EMBL Data Library, December 1992 A;Description: Promoter first exon/intron characte A;Reference number: 843170 A;Reference number: 843170 A;Reference number: DNA A;Residues: 1-12 <che> A;Cross-references: UNIPROT:Q13865; EMBL:X69658; N</che>
	0000000000000000000	fra re bra ex
	11111 12222 12222	- human (fragment lens (man) #sequence_revision Brown, A.; Jones, 3L Data Library, Duter first exon/in/ \$43170
	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	nain - h sapiens 1995 #se 1170 Y.; Brc Y.; Brc Y.; Brc Y.; Brc Y.; Brc Ininary
	U U U U U U U U U U U U U U U U U U U	light chain is: Homo sapi 13-Jan-1995 iion: S43170 ijovsky, Y.; id to the EME ption: Promc proe number: iion: S43170 ii: preliminar iie type: DNF
	и м м м м м м м м м м м м м м м м м м м	RESULT 1 S43170 kinesin light chain - human (fragmer C; Species: Homo sapiens (man) C; Date: 13-Jan-1995 #sequence_revisiC; Accession: S43170 R; Chernajovsky, Y.; Brown, A.; Jones submitted to the EMBL Data Library, A; Description: Promoter first exon/iA; Reference number: S43170 A; Accession: S43170 A; Accession: S43170 A; Accession: Lype: DNA A; Residues: 1-12 < CHE> A; Cross-references: UNIPROT:Q13865;

.; 0 ö Belong

Big H2 chain - Pacific hagfish (fragment)

C; Species: Eptatretus stouti (Pacific hagfish)

C; Species: Bptatretus stouti (Pacific hagfish)

C; Date: 03-Jun-1993 #sequence_revision 03-Jun-1993 #text_change 03-Jun-1993

C; Accession: B60769

R; Hanley, P.J.; Seppelt, I.M.; Gooley, A.A.; Hook, J.W.; Raison, R.L.

J. Immunol. 145, 3823-3828, 1990

A; Title: Distinct Ig H chains in a primitive vertebrate, Eptatretus stouti.

A; Reference number: A60769; MUID:91060965; PMID:2123225

A; Accession: B60769

A; Accession: preliminary

A; Molecule type: protein

A; Residues: 1-11 < HAN> Gaps Gaps .. ., 47.5%; Score 19; DB 2; Length 11; 80.0%; Pred. No. 5.6e+02; Live 0; Mismatches 1; Indels Length 12 50.0%; Score 20; DB 2; I 60.0%; Pred. No. 3.8e+02; iive 2; Mismatches 0; Conservative 4; Conservative Best Local Similarity Matches 3, Conserv Query Match Best Local Similarity 3 MIQGG 7 1 MLRGG 5 Query Match Matches RESULT 2 g ð

1 DXWIQ 5 DEMIQ 9 RESULT 3 PT0542 à

T-cell receptor beta chain V-D-J region (126-1BA) - mouse (fragment)

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C;Accession: S60354
R;Huang, D.Y.; Ichikawa, Y.
Biochim. Biophys. Acta 1243, 431-436, 1995
A;Title: Identification of essential lysyl and cysteinyl residues, and the amino acid sec A;Reference number: S60354; MUID:95244596; PMID:7727518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kawasaki Igakkaishi 22, 245-259, 1996
A;Title: Analysis of proteins isolated by two dimensional electrophoresis of mouse neurok
A;Reference number: PN0041
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Molecule type: protein
A;Residues: 1-11 <XAT>
A;Experimental source: neuroblastoma cell
C;Comment: The molecular mass is 13,900 and the pI is 6.36. The amino-terminus is blockec
C;Keywords: brain
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C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Species: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change 30-May-1997
C;Accession: PT0218
R;Nakano, N.; Kikutani, H.; Nishimoto, H.; Kishimoto, T.
J. Exp. Med. 173, 1091-1097, 1991
A;Title: T cell receptor V gene usage of islet beta cell-reactive T cells is not restrict A;Reference number: PT0209; MUID:91217621; PMID:1902501
A;Reference number: PT0209; MUID:91217621; PMID:1902501
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein kinase C inhibitor I - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 29-Oct-1997 #sequence_revision 29-Oct-1997 #text_change 23-Jan-1998
C;Accession: PN0044
                                                                                 C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 24-Aug-1996 #sequence_revision 13-Mar-1997 #text_change 13-Mar-1997
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Best Local Similarity 100.0%; Pred. No. 9.2e+02;
Matches 3; Conservative 0; Mismatches 0;
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75.0%; Pred. No. 9.2e+02;
tive 1; Mismatches 0;
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                                                              retinal oxidase - rabbit (fragment)
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Best Local Similarity 100.
Matches 3; Conservative
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A, Residues: 1-11 <NAK>
C, Keywords: T-cell receptor
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Best Local Similarity
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A;Molecule type: protein
A;Residues: 1-11 <HUA>
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PH1343
Ig heavy chain DJ region (clone C100-91) - human (fragment)
C; Species: Howo sapiens (man)
C; Species: Howo sapiens (man)
C; Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C; Accession: PH1343
R; Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
J. Exp. Med. 176, 1577-1581, 1992
A; Title: Predominance of fetal type DJH joining in young children with B precursor lymph A; Reference number: PH1302; MUID: 93094761; PMID: 1460419
A; Reference number: PH1343
A; Molecule type: DNA
A; Residues: 1-11 < WAS>
C; Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Molecule type: protein
A;Residues: 1-9 <MON>
A;Cross-references: UNIPROT:P01158
C;Comment: This peptide was obtained from dialysates of occipital venous sinus blood fro
of recipient rabbits, it induces spindle and delta EEG activity and reduced motor activ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            delta sleep-inducing peptide - rabbit
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 20-Jun-2000 #sequence_revision 20-Jun-2000 #text_change 16-Aug-2004
C;Accession: A01422
R;Monnier, M.; Dudler, L.; Gachter, R.; Maier, P.F.; Tobler, H.J.; Schoenenberger, G.A.
Experientia 33, 548-552, 1977
A;Title: The delta sleep inducing peptide (DSIP). Comparative properties of the original
A;Reference number: A01422; MJID:77185324; PMID:862769
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0542
R;Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Accession: PT0542
A;Status: translation not shown
A;Molecule type: mRNA
A;Residues: 1-7 <FEE>
A;Experimental source: day 18 fetal thymus, strain BALB/C
C;Keywords: T-cell receptor
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C;Species: Sus scrofa domestica (domestic pig)
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 13-Jul-1981 #sequence_revision 13-Jul-1981 #text_change 09-Jul-2004
C;Accession: A01523; A60983
R;Pleau, J.M.; Dardenne, M.; Blouquit, Y.; Bach, J.F.
J. Biol. Chem. 252, 8045-8047, 1977
A;Title: Structural study of circulating thymic factor: a peptide isolated from pig serun A;Reference number: A01523; MUID:78026571; PMID:914862
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C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
C;Accession: A60957
R;Ernstroem, U.; Gafvelin, G.; Rudja, J.M.
Biosci. Rep. 10, 403-412, 1990
A;Title: Purification of thymocyte growth peptide (TGP) from sheep thymus. Relationship t
A;Reference number: A60957; MUID:91064427; PMID:2249004
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A; Residues: 19 < ERN>
A; Residues: 19 < ERN>
A; Cross-references: UNIPROT: Q7M3C5
C; Comment: This peptide induces DNA synthesis in immature thymocytes, but not peripheral in a variety of immunoassays.
C; Comment: This peptide was isolated in two forms. One form contained the pyrrolidone can r form (thymocyte growth peptide) contains a large, non-peptide blocking group with a high sympactamily: thymic factor
C; Superfamily: thymic factor
C; Keywords: blocked amino end; pyroglutamic acid
C; Keywords: blocked amino end (Glx) (in FTS) #status experimental
F; I/Modified site: blocked amino end (Glx) (in thymocyte growth peptide) #status experime
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Best Local Similarity 100.
Matches 3; Conservative
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RESULT 12
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N;Alternate names: hemorrhagic toxin
C;Species: Trimeresurus elegans (Sakishima habu)
C;Date: 20-Oct-1994 #sequence_revision 06-Jan-1995 #text_change 06-Jan-1995
C;Accession: A61575
R;Nikai, T.; Komori, Y.; Imai, K.; Sugihara, H.
Int. J. Biochem. 23, 73-78, 1991
A;Title: Isolation and characterization of hemorrhagic toxin from the venom of Trimeresu
A;Reference number: A61575; MUID:91216327; PMID:2022298
                                                          T-cell receptor gamma chain (5t.1) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 07-May-1999
C;Accession: 141946
R;Whetsell, M.; Mosley, R.L.; Whetsell, L.; Schaefer, F.V.; Miller, K.S.; Klein, J.R.
Mol. Cell. Biol. 11, 5902-5909, 1991
A;Title: Rearrangement and junctional-site sequence analyses of T-cell receptor gamma 2A;Reference number: A41946; MUID:92049316; PMID:1658619
A;Accession: 141946
A;Accession: 141946
A;Status: preliminary; not compared with conceptual translation
A;Molecule type: DNA
A;Residues: 1-11 < WHE>
C;Keywords: T-cell receptor
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R.HSi, K.L.; Chen, R.L.; Chen, Z.G.; Zhang, H.L.; Lu, Y.A.; Guo, S.Y.; Wu, S.X.; Tsou, Arch. Biochem. Biophys. 240, 178-183, 1985
A;Reference number: A23751; MUID:85250425; PMID:4015098
A;Accession: A23751
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-3 <HSI>
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C;Species: Sus scrofa domestica (domestic pig)
C;Date: 15-Jun-2001 #sequence_revision 15-Jun-2001 #text_change 15-Mar-2004
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Pred. No. 9.2e+02;
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100.0%; Pred. No. 9.2e+02;
iive 0; Mismatches 0; Indels
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100.0%; Pred. No. ..
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A;Molecule type: protein
A;Residues: 1-11 <NIK>
C;Keywords: hydrolase; serine proteinase; venom
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RESULT 15
D37196
bradykinin-potentiating peptide 4 - island jararaca
C; Species: Bothrops insularis (island jararaca)
C; Date: 14-Fb-1992 #sequence_revision 01-Dec-1992 #text_change 09-Jul-2004
C; Accession: D37196
R; Cintra, A.C.O.; Vieira, C.A.; Giglio, J.R.
J. Protein Chem. 9, 221-227, 1990
A; Title: Primary structure and biological activity of bradykinin potentiating peptides facence number: A37196; MUID:90351557; PMID:2386615
A; Accession: D37196
A; Status: preliminary
A; Molecule type: protein
A; Residues: 1-11 < CIN>
A; CIN>
A; Conserver experiments
C; Keywords: pyroglutamic acid
F; 1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
S57575
T cell receptor V-J junctional alpha chain region - human (fragment)
C; Species: Homo sapiens (man)
C; Date: 19-Oct-1995 #sequence_revision 17-Nov-1995 #text_change 05-Nov-1999
C; Accession: S57575
R; Burrows, S.R.; Silins, S.L.; Moss, D.J.; Khanna, R.; Misko, I.S.; Argaet, V.P.
submitted to the EMBL Data Library, June 1995
A; Description: T cell receptor repertoire for a viral epitope in humans is diversified A; Reference number: S57494
A; Recession: S57575
A; Ratus: preliminary
A; Molecule type: mRNA
A; Residues: 1-11 < BUR>
A; Residues: 1-11 < BUR>
A; Cross-references: EMBL: Z49953; NID: G887510; PIDN: CAA90224.1; PID: G887511
C; Keywords: T-cell receptor
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Search completed: October 21, 2004, 07:27:58 Job time : 16.75 secs

OGG 3

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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- protein search, using sw model OM protein

October 21, 2004, 07:05:04; Search time 86 Seconds (without alignments) 60.214 Million cell updates/sec Run on:

US-09-720-469A-38 40 1 DXMIQGGDX 9

Perfect score:

Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

1825181 segs, 575374646 residues Searched:

4909 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 12

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

UniProt_02:* Database

1: uniprot_sprot:*
2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	Q63480 rattus norv	Q9h3w2 homo sapien	Q70y58 clinopodium		Q13865 homo sapien	rattu	Q70y98 capitanopsi		P01158 oryctolagus	Q61dh6 rattus norv	0 rattu	Q51812 plasmid f.	homo sa				Bacc	P84071 allium asca	P01255 sus scrofa	Q7m3c5 ovis aries	homo	Q96qa7 homo sapien	Aad14854 homo sapi				2	P81012 schizaphis	a	Q7m0a7 cricetidae	Q8uvv8 squalus aca
SUMMARIES	ID	Q63480	Q9H3W2	Q70Y58	CAD45546	Q13865	055184	Q70Y98	CAD45456	DSIP_RABIT	<u> Фегрн</u> е	AAA40600	051812	921H6Q	O9N2B8	Q9N2B9	Q9N2C0	054226	ASCL_ALLAS	THYF_PIG	Q7M3C5	Q6LC14	Q96QA7	AAD14854	BPP4_BOTIN	Q70Y64	Q47420	CAD45529	ESTA_SCHGA	QBUVW2	Q7M0A7	QBUVVB
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o <b>k</b> c	Query Match	7.	δ.	ς,	•	50.0	•	-	7.		45.0	45.0	45.0	45.0					42.5			•						ς.	•	0	40.0	40.0
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;

[1]
SEQUENCE FROM N.A.
MEDLINE=94010310; PubMed=8406013;
Onno M., Nakamura T., Hillova J., Hill M.;

Created)
Last sequence update)
Last annotation update)

Q9H3W2; 01-MAR-2001 (TrEMBLrel. 16, C: 01-MAR-2001 (TrEMBLrel. 16, L: 01-JUN-2003 (TrEMBLrel. 24, L: TRE17 protein. Name=TRE17 gene; Homo sapiens (Human).

12 AA

PRT;

PRELIMINARY;

Q9H3W2

RESULT 2 Q9H3W2

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Q8uvw0 rana catesb	P18691 thunnus alb	Q9p0k3 homo sapien	Q9bgt4 homo sapien	Q6y£34 rattus norv	Q92009 gallus gall	Aan87279 rattus no	Q6x2s9 homo sapien	Q85v66 eucalyptus	Q71zj7 carcharhinu	Q7zzj2 motacilla f	Aap83315 homo sapi	Q7zzi6 motacilla f	Q7zzi9 motacilla f
OBUVWO	ACI THUAL	<u>оэрокз</u>	Q9BQT4	Q6YF34	600560	AAN87279	Q6 <b>X</b> 2S9	Q85V66	Q7LZJ7	Q7ZZJ2	AAP83315	Q7ZZ16	Q7ZZI9
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## ALIGNMENTS

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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Splice variants of rat TR4 orphan receptor: differential expression of novel sequences in the 5'-untranslated region and C-terminal
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MEDLINE=96198747; PubMed=8612486;
Yoshikawa T., Makino S., Gao X.M., Xing G.Q., Chuang D.M.,
Detera-Wadleigh S.D.;
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                                                                                                                             Last sequence update)
Last annotation update)
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7 AA.
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EMBL; U59125; AAB02827.1; -.
GO; GO:0004872; F:receptor activity; IEA.
                                                                              01-NOV-1996 (TrEMBLrel. 01, Created) 01-NOV-1996 (TrEMBLrel. 01, Last seq 01-JUN-2003 (TrEMBLrel. 24, Last ann TR4-NS orphan receptor (Fragment).
   PRT;
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PRELIMINARY;
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1 MLRGG
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EMBL; AJ505426; CAD45546.1; -. GO: GO:0003735; F:structural constituent of ribosome; IEA.
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
lamiids; Lamiales; Lamiaceae; Nepetoideae; Nepeteae; Clinopodium.
"Identification of novel sequences in the repertoire of hypervariable TRE17 genes from immortalized nonmalignant and malignant human keratinocytes.";
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                                                                                                        SEQUENCE FROM N.A.
MEDLINE=93228825; PubMed=8471161;
Onno M., Nakamura T., Mariage-Samson R., Hillova J., Hill M.;
"Human TRE17 oncogene is generated from a family of homologous polymorphic sequences by single-base changes.";
DNA Cell Biol. 12:107-118(1993).
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                                                                                                                                                                                                                                                                       Hillova F.;
Submitted (APR-1993) to the EMBL/GenBank/DDBJ databases.
EMBL; X71371; CAC16150.1; -.
SEQUENCE 12 AA; 1252 MW; 6E17AFDE83ADD87B CRC64;
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
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100.0%; Fic
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                                                                 Gene 131:209-215(1993)
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Best Local Similarity
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CAD45546
ID CAD4
AC CAD4
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DT 14-M
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EMBL; AJ505426; CAD45546.1; -. Chloroplast; Ribosomal protein.

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SEQUENCE 12 AA; 1353 MW; D43D1B51AF587866 CRC64;
                           Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Varidiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; Core eudicots; asterids; lamiales; Lamiaceae; Nepetoideae; Nepeteae; Clinopodium; Clinopodium vulgare.
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MEDLINE=97101110; PubMed=8945637;
Chernajovsky Y., Brown A., Jones T.A.;
Chernajovsky Y., Brown A., Jones T.A.;
"Promoter first exon/intron characterization and chromosomal location of the human light chain (beta) kinesin gene.";
DNA Cell Biol. 15:965-974(1996).
EMBL; X69658; CAA49349.1; -.
PIR; S43170; S43170.
SEQUENCE 12 AA; 1274 MW; 3FB8D34EE165A5B8 CRC64;
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
NCBI_TaxID=9606;
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055184;
01-JUN-1998 (TrEMBLrel. 06, Created)
01-JUN-1998 (TrEMBLrel. 06, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
0rphan receptor TR4-NS (Fragment).
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Last annotation update)
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60.0%; Pred. No. 3.5e+03;
iive 2; Mismatches 0;
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100.0%; Pred. No. 2.1e+03;
tive 0; Mismatches 0;
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Clinopodium vulgare subsp. arundanum.
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01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last seq
01-JUN-2003 (TrEMBLrel. 24, Last anno
Kinesin light chain.
Name-beta-kinesin;
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Capitanopsis angustifolia.
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                                                                                                                                               PRELIMINARY;
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Best Local Similarity
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1 LFQGG
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P01158;
21-JUL-1986
21-JUL-1986
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CAD45456
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
lamiids; Lamiales; Lamiaceae; Nepetoideae; Ocimeae; Capitanopsis.
NCBL_TaxID=204110;
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EMBL, AJ505333; CAD45456.1; -. GO: GO:0003735; F:structural constituent of ribosome; IEA.
                                            Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
NCBI_TaxID=10116;
                                                                                                                                                                   STRAIN=Sprague-Dawley;
MEDLINE=96198747; PubMed=8612486;
Yoshikawa T., Makino S., Gao X.M., Xing G.Q., Chuang D.M.,
Detera-Wadleigh S.D.;
"Splice variants of rat TR4 orphan receptor: differential expression of novel sequences in the 5'-untranslated region and C-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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EMBL; US9454; AAB91433.1; -.
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80.0%; Pred. No. 1.8e+06;
tive 0; Mismatches 1; Indels
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN=Sprague-Dawley;
MEDLINE=96299786; PubMed=8661150;
                                                                                                                                                                                                                                                                                                                                          Endocrinology 137:1562-1571(1996)
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Name=rps16;
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Best Local Similarity 60.0
Matches 3; Conservative
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                             Rattus norvegicus (Rat)
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                                                                                                                                                                                                                                                                                                                     domain.";
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3 MIQGG 7

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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
lamiids; Lamiales; Lamiaceae; Nepetoideae; Ocimeae; Capitanopsis.
NCBI_TaxID=204110;
                                                                                                                                                                                                                                                                                                                                      Paton A. Springate D.A., Sudde S., Otieno D., Grayer R., Harley M.M., Willis F., Simonds M.S.J., Powell M.P., Savolainen V.; "Phylogeny and evolution of basils and allies (Ocimeae, Labiatae) based on three plastid DNA regions."; Mol. Phylogenet. Evol. 31:277-299(2004).
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MEDLINE=79054421; PubMed=568769;
Schoenenberger G.A., Maier P.F., Tobler H.J., Wilson K., Monnier M.;
"The delta EEG (sleep)-inducing peptide (DSIP). XI. Amino-acid analysis, sequence, synthesis and activity of the nonapeptide.";
Pflugers Arch. 376:119-129(1978).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "The delta sleep inducing peptide (DSIP). Comparative properties of
the original and synthetic nonapeptide.";
Experientia 33:548-552(1977).
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Monnier M., Dudler L., Gachter R., Maier P.F., Tobler H.J.,
Schoenenberger G.A.;
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Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.

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    SEQUENCE
    11 AA;
    1161 MW;
    2AB3C5D9BEAB5878
    CRC64;

                                       14-MAR-2004 (TrEMBLrel. 27, Created)
14-MAR-2004 (TrEMBLrel. 27, Last sequence update)
14-MAR-2004 (TrEMBLrel. 27, Last annotation update)
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Graf M.V., Kastin A.J.;
"Delta-sleep-inducing peptide (DSIP): an update.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Delta sleep-inducing peptide (DSIP).
Oryctolagus cuniculus (Rabbit).
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11 AA
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Local Similarity
les 3; Conserv
NCBI_TaxID=10116;
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NCBI_TaxID=2465;
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Plasmid F.
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SEQÜENCE
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Q9H1Z6;
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MEDLINE=92129344; PubMed=1733955;
Krozowski Z., Obeyesekere V., Smith R., Mercer W.;
"Tissue-specific expression of an 11-beta-hydroxysteroid dehydrogenase with a truncated N-terminal domain: A potential mechanism for differential intracellular localization within mineralocorticoid
                                                                                  reduced motor activities.
-!- MISCELLANEOUS: This peptide was obtained from dialysates of occipital venous sinus blood from rabbits kept asleep by electric stimulation of the thalamus.
-!- DATABASE: NAME=Protein Spotlight; NOTE=Issue 8 of March 2001; WWW="http://www.expasy.org/spotlight/articles/sptlt008.html".
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Rattus.
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EMBL; M77835; AAA40600.1; -.

GO; GO:0003845; F:11-beta-hydroxysteroid dehydrogenase activity; IEA.

GO; GO:0016491; F:oxidoreductase activity; IEA.
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                                -1- FUNCTION: When infused into the mesodiencephalic ventricle of recipient rabbits induces spindle and delta EBG activity and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
11-beta-hydroxysteroid dehydrogenase (EC 1.1.1.146) (Fragment).
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05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
11-beta-hydroxysteroid dehydrogenase (EC 1.1.1.146) (Fragment)
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9 AA; 1015 MW; 41F942C3333866D7 CRC64;
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Direct protein sequencing.
SEQUENCE 9 AA; 849 MW; DDD365BDDAA8787D CRC64;
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Pred. No. 1.8e+06;
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75.0%;
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02-MAR-2004 (TrEMBLrel. 27,
02-MAR-2004 (TrEMBLrel. 27,
02-MAR-2004 (TrEMBLrel. 27,
    Peptides 7:1165-1187(1986)
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Best Local Similarity lov.
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nes 3; Conservative
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J. Biol. Chem. 26
EMBL; M77835; AAA
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QGLDHG
ID QGLDH
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DT 05-JU
DE 11-be
GN Name=
OS Rattu
OC Mamma
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                                             Krozowski Z., Obeyesekere V., Smith R., Mercer W.;
"Tissue-specific expression of an 11-beta-hydroxysteroid dehydrogenase with a truncated N-terminal domain: A potential mechanism for differential intracellular localization within mineralocorticoid target cells.";
J. Biol. Chem. 267:2569-2574 (1992).
EMBL; M77835; AAA40600.1; -.
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                                                                                                                                                                                                                                                                                                                                                                               Gaps
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                  Query Match 45.0%; Score 18; DB 2; Length 9; Best Local Similarity 75.0%; Pred. No. 1.8e+06; Matches 3; Conservative 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                    41F942C3333866D7 CRC64;
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Last annotation update)
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Last annotation update)
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GO; GO:0004386; F:helicase activity; IEA.
Helicase; Plasmid.
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STRAIN=Sprague-Dawley;
MEDLINE=92129344; PubMed=1733955;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A. MEDLINE-92009201; PubMed=1916281;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         051812;
01-NOV-1996 (TrEMBLrel. 01,
01-DEC-2001 (TrEMBLrel. 19,
01-JUN-2003 (TrEMBLrel. 24,
DNA helicase (Fragment).
                                                                                                                                                                                                                                                                  SEQUENCE 9 AA; 1015 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRELIMINARY;
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Search completed: October 21, 2004, 07:26:49 Job time: 88 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                Pongo pygmaeus (Orangutan).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Pongo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gorilla górilla (gorilla).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Gorilla.
NCBI_TaxID=9593;
                                                                                                                                                                                             .;
0
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                                                                                                                                                            Score 18; DB 2; Length 12;
Pred. No. 9.2e+03;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         45.0%; Score 18; DB 2; Length 12; 100.0%; Pred. No. 9.2e+03; ative 0; Mismatches 0; Indels
          Gonzalez-Gonzara F., Pedrosa C., Ramirez J.P.,
Castilla J.A., Yoldi A., Ruiz A., Real L.M.;
Submitted (DEC-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF326912; AAG42501.1; -.
GO; GO:0004872; F:receptor activity; IEA.
Receptor. 1 1
NON_TER 12 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
Kitano T., Kobayakawa H., Saitou N.;
Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases.
EMBL, AB041365; BAA94454.1; -.
GO; GO:0008168; F:methyltransferase activity; IEA.
GO; GO:0016740; F:transferase activity; IEA.
Methyltransferase; Transferase.
NON TER
12
12
SEQUENCE 12 AA; 1290 MW; CF079554917861A9 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
Kitano T., Kobayakawa H., Saitou N.;
Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AB041364; BAA94453.1; -.
GO; GO:0008168; F:methyltransferase activity; IEA.
GO; GO:0016740; F:transferase activity; IEA.
                                                                                                    1 1
12 12
12 AA, 1385 MW, 09168BCB76C41404 CRC64;
                                                                                                                                                                                                                                                                                                                                         09N2BB;
01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Indolethylamine N-methyltransferase (Fragment).
Name=INMT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Indolethylamine N-methyltransferase (Fragment).
                                                                                                                                                                                                                                                                                                                               12 AA.
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                                                                                                                                                           Best Local Similarity 57.18;
Matches 4; Conservative (
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Best Local Similarity
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SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9600;
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Q9N2B9;
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                                                                Length 12;
                                                                                              0; Indels
Methyltransferase; Transferase.

NON TER 12 12

SEQUENCE 12 AA; 1289 MW; CF0798E2917861A9 CRC64;
                                                              Similarity 100.0%; Pred. No. 9.2e+03; 3; Conservative 0; Mismatches 0;
                                                               Query Match
Best Local Similarity
                                                                                                                            6 GGD 8
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 us-09-720-469a-39. rag

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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- protein search, using sw model OM protein Run on:

October 21, 2004, 07:04:19; Search time 83 Seconds (without alignments) 38.898 Million cell updates/sec

US-09-720-469A-39 50 1 KYHRVIKDF 9 Title: Perfect score:

Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

2002273 seqs, 358729299 residues Searched:

497672 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 12

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

A_Geneseq_23Sep04:* 1: genesernleer Database :

geneseqp1980s:*
geneseqp1990s:*
geneseqp2000s:*
geneseqp2001s:*
geneseqp2003as:*
geneseqp2003as:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	Aav69962 Human cvc	Aab46946 Human cvc	1 Human	0 Human	Antitu	Abg79075 Human Cyp	Human	Adc17701 Cyclophiî	Adg89641 Class I H	Aay69960 Human cyc	1 Hepat:	Aab82292 Phosphory	Aay69959 Human cyc	₩	Н		Н	Abu57322 Replikin	Add88394 Tumour vi	Add88395 Tumour vi	Adg18172 Tumour vi	Adg18173 Tumour vi	7	Aau71498 Human MHC	Aan71218 Human MHC
SUMMARIES		AAY69962	AAB46946	AAY69921	AAB46930	AAG68088	ABG79075	ABR84373	ADC17701	ADG89641	AAY69960	AAR96521	AAB82292	AAY69959	AAY69924	ABG94771	ABG94772	ABU57321	ABU57322	ADD88394	ADD88395	ADG18172	ADG18173	ADI46367	AAU71498	AAU71218
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Gaps

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Aau71556 Human MHC	Aau71745 Human MHC	Adm41409 Human bet	Aaw92359 CYP2Bl im	Aar48203 Human pho	Aab82291 Phosphory	Aab82294 Phosphory	Aap93365 Amino aci	Abp11571 HIV A01 s	Abp15315 HIV A24 s	Abp20150 HIV A03 m	Abp15417 HIV A24 s	Abp11625 HIV A01 s	Abp15505 HIV A24 s	Abp20151 HIV A03 m	Abp17665 HIV B58 s	Abp15503 HIV A24 s	Adb79463 Parapoxvi	Adp74879 Parapoxvi	7
AAU71556	AAU71745	ADM41409	AAW92359	AAR48203	AAB82291	AAB82294	AAP93365	ABP11571	ABP15315	ABP20150	ABP15417	ABP11625	ABP15505	ABP20151	ABP17665	ABP15503	ADB79463	ADP74879	ABR11047
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10	10	10	12	11	11	11	12	σ	σ	10	10	11	11	11	11	11	7	7	σı
54.0	54.0	54.0	54.0	52.0		52.0	52.0	51.0	51.0	51.0	51.0	51.0	51.0	51.0	51.0	51.0	50.0	50.0	50.0
27			27		26	26	26	25.5		25.5	25.5	25.5	25.5	25.5	S	'n.		25	
26	27	28	29	30	31	32	33	34		36	37	38	39	40	41	42	43	44	45

### ALIGNMENTS

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This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                 Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                           Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 50; DB 3; Length 9; ilarity 100.0%; Pred. No. 1.7e+06; Conservative 0; Mismatches 0; Indels
                                                                                          Human cyclophilin B peptide fragment #42.
                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 10; Page 60; 64pp; Japanese.
              AAY69962 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                    (SUMU ) SUMITOMO PHARM CO LTD.
                                                                                                                                                                                                                                  99WO-JP003360.
                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                WPI; 2000-116932/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                         Gomi S;
                                                                                                                                                                                                                                                                                                (ITOH/) ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9 AA;
                                                                                                                                                                                WO9967288-A1.
                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                   24-JUN-1999;
                                                                                                                                                                                                          29-DEC-1999.
                                                                                                                                                                                                                                                           25-JUN-1998;
                                                                11-APR-2000
                                       AAY69962;
                                                                                                                                                                                                                                                                                                                        Itoh K,
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1 KYHRVIKDF

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This invention describes the novel use of interferons (IFNS) or DNAs capable of expressing the interferons and/or antigenic proteins (AP), antigenic peptides derived from the proteins or DNAs capable of expressing the antigeners or peptides, in the manufacture of an agent for induction of antigen-specific T cells. The products of the invention have virucide and cytostatic activity and can be used for gene therapy or as inducers of antigen-specific T cells. The action of interferon-alpha (IFN-alpha) in a system for inducing specific cytotoxic T cell (CTL) by administering an antigenic peptide in an incomplete Freund's adjuvant (IFNA) emulsion preparation form was evaluated. IFNS (DNA encoding IFNS) are useful in the manufacture of a medicament for inducing antigen-specific T cells in an individual who has been administered with AP (or DNA encoding AP) or vice versa. The medicament is useful for the treatment or prophylaxis of a tumor or a viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Use of antigenic proteins, peptides, interferon or their encoding DNA, in the manufacture of an agent for the induction of antigen-specific {\tt T}
                                                                                                                                                            Tumor antigenic peptide; interferon; IFN; antigen-specific T cell; virucide; cytostatic; gene therapy; IFN-alpha; cytotoxic T cell; viral infectious disease; cyclophilin B; human.
                                                                                                                               Human cyclophilin B derived tumor antigenic peptide SEQ ID 32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 50; DB 4; Length 9; 100.0%; Pred. No. 1.7e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY69921 standard; peptide; 9 AA.
                               AAB46946 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Yamaoka T;
                                                                                                                                                                                                                                                                                                                                                                                                  (SUMU ) SUMITOMO PHARM CO LID
                                                                                                                                                                                                                                                                                                                                 24-JUL-2000; 2000EP-00306263.
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                                                                                                (first entry)
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                                                                                                                                                                                                                                                                EP1074267-A1
                                                                                                04-MAY-2001
                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                  22-JUL-1999;
                                                                                                                                                                                                                                                                                                07-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY69921;
                                                                AAB46946;
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AC AAX6
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RESULT 2
AAB46946
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Gaps

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This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                             Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                       Tumor antigen peptides derived from cyclophilin E for treatment and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tumor antigenic peptide; interferon; IFN; antigen-specific T cell; virucide; cytostatic; gene therapy; IFN-alpha; cytotoxic T cell; viral infectious disease; cyclophilin B; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human cyclophilin B derived tumor antigenic peptide SEQ ID 16.
                                                                                                                                                                                                                                                                                                                                                                                                                   92.0%; Score 46; DB 3; Length 9; 88.9%; Pred. No. 1.7e+06; ive 1; Mismatches 0; Indels
         Human cyclophilin B peptide fragment #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB46930 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                            Claim 4; Page 49; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Takasu H, Gotoh M, Yamaoka T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SUMU ) SUMITOMO PHARM CO LTD.
                                                                                                                                                                                        (SUMU ) SUMITOMO PHARM CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JUL-2000; 2000EP-00306263.
                                                                                                                                         99WO-JP003360.
                                                                                                                                                                98JP-00178449.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                     diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-193144/20
                                                                                                                                                                                                                                                WPI; 2000-116932/10.
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 KYHRVIKDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KFHRVIKDF
                                                                                                                                                                                                                           Gomi S;
                                                                                                                                                                                                    ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                             Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-JUL-1999;
                                                                                                                                          24-JUN-1999;
                                                                                                                                                                 25-JUN-1998;
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                                                                     Homo sapiens
                                                                                           WO9967288-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-MAY-2001
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                                                                                                                  29-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB46930;
                                                                                                                                                                                                                          Itoh K,
                                                                                                                                                                                                     (ITOH/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 4
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Use of antigenic proteins, peptides, interferon or their encoding DNA, in the manufacture of an agent for the induction of antigen-specific {\tt T}
                                                                                                                                                                                                                                                                           This invention describes the novel use of interferons (IFNs) or
                                                                                                                                                                                      Disclosure; Page 15; 25pp; English
                                                                                             cells
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capable of expressing the interferons and/or antigenic proteins (AP), antigenic peptides derived from the proteins or DNAs capable of expressing the antigenic proteins or peptides, in the manufacture of an agent for induction of antigen-specific T cells. The products of the invention have virucide and cytostatic activity and can be used for gene therapy or as inducers of antigen-specific T cells. The action of interferon-alpha (IFN-alpha) in a system for inducing specific cytotoxic T cell (CTL) by administering an antigenic peptide in an incomplete Freund's adjuvant (IFN) emulsion preparation form was evaluated. IFNs (or inducing antigen-specific T cells in an individual who has been administered with AP (or DNA encoding AP) or vice versa. The medicament is userious discretions discretions or a viral infectious disease Sequence 9 AA; 

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                             Gaps
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          Score 46; DB 4; Length 9;
Pred. No. 1.7e+06;
1; Mismatches 0; Indels
           92.0%;
88.9%;
Query Match
Best Local Similarity 86.,
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6 1 KYHRVIKDF |:|||||| 1 KFHRVIKDF

AAG68088 standard; peptide; 9 AA. 17-DEC-2001 AAG68088; RESULT 5 

Antitumour peptide cyclophilin B 84-92 (first entry)

Antitumour; cancer; cancer cell recognition; antigenic; CTL; lck; src; tumour specific cytotoxic T lymphocyte; anticancer; SART-1; SART-3; cyclophilin B gene; HLA-A2402.

Homo sapiens

JP2001245675-A.

11-SEP-2001

25-DEC-2000; 2000JP-00393047.

99JP-00374322 28~DEC-1999;

Y OTI (/YOTI)

WPI; 2001-610076/70.

New peptides for recognizing cancer cells with tumor specific cytotoxic T lymphocytes and for treating cancer.

Claim 8; Page 2; 14pp; Japanese.

The present invention describes peptides recognising cancer cells with tumour specific cytotoxic T lymphocytes (CTL). The peptides recognising cancer cells with tumour specific CTLs are selected from: (1) peptides of sequences (AAG68066 to AAG68069); (2) peptides containing the above mentioned sequences; (3) peptides having 70 % or more of homogeneity with the above mentioned sequences; and (4) peptides with one or more deleted,

The invention relates to a composition (I) comprising an immune effector cell and a cell penetrating peptide (CPP) associated with an antigen or antibody. Also included are (1) a vaccine comprising (I), CPP associated with an antigen, and a pharmaceutically acceptable carrier and (2) preparing a composition for a disease, by providing (I) and CPP associated with an antigen for disease, and introducing the antigenassociated CPP to (I), where antigen enters into the cell. The antigenare, for example, tumour antigen derived epitopes recognised by tumour infiltrating lymphocytes (TIL) of HLA (human leukocyte antigen) class I or II. The composition is useful for enhancing immunity in an animal to a disease, by administering a mature dendritic cell comprising CPP associated with an antigen to disease, to the animal, such that following

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            sequences, particularly those having recognising property due to HLA-A2402 binding CTL, especially having at least 5 amino acids, used for medicine, particularly anticancer agents, derived from antitumour antigenic peptides of lck, src family, SART-1, SART-3 or cyclophilin B genes. The antitumour peptides have cytostatic activities. The peptides are used for the treatment of cancer. The peptides cause activation of CTL in cancer patients. The present sequence represents a peptide from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         penetrating peptide; cancer; tumour; melanoma; thymoma; antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    lymphoma; sarcoma; lung cancer; non-Hodgkin's lymphoma; leukaemia; Hodgkin's lymphoma; leukaemia; Hodgkin's lymphoma; uterine cancer; cervical cancer; bladder cancer; kidney cancer; adenocarcinoma; breast cancer; prostate cancer; ovarian cancer; pancreatic cancer; epitope; vaccine; dendritic cell; tumour infiltrating lymphocyte; IIL; human leukocyte antigen; HLA;
                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel composition for treating a disease in an animal, comprises an immune effector cell and cell penetrating peptide associated with an antigen or antibody.
substituted, added or inserted amino acid(s) of the above mentioned
                                                                                                                                                                                                                .,
                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Cyp-B class I HLA widely expressed antigen peptide #1.
                                                                                                                                                                  Score 46; DB 4; Length 5, Pred. No. 1.7e+06;
                                                                                                                                                                                                               1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 17; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                   ABG79075 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (BAYU ) BAYLOR COLLEGE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-FEB-2002; 2002WO-US005212.
                                                                                                                                                                                92.0%;
88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-FEB-2001; 2001US-0268687P.
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                                                                                                                                                                                                               8; Conservative
                                                                                                                       the present invention
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                                                                                                                                                                                               Best Local Similarity
                                                                                                                                                                                                                                           1 KYHRVIKDF
                                                                                                                                                                                                                                                                       1 KFHRVIKDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cytostatic; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200264057-A2
                                                                                                                                                  Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-AUG-2002
                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wang R;
                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                 RESULT 6
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Sequence 9 AA;

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The invention relates to a method for the detection of antigen specific T -cells in a blood sample involving the use of a plurality of antigenic peptides. The method comprises sampling of peripheral blood monocytes; stimulation of the collected peripheral blood monocytes with antigens without direct use of antigen presenting cells; and detection of T-cells specific to the antigen in the stimulated monocytes. The method is particularly used for the detection of cancer as it can be used in semicantitative determination of cancer specific T-cells. It can also be used for cancer vaccine therapy for patients with cervical or prostate cancer. The method can additionally be used to monitor of cellular immunity and cancer immune therapy by detection of specific T-cell frequencies. Sequences ABR84367-ABR84380 represent HLA-A24 (human leukocyte antigen) peptides of human origin used in an example from the invention
                                                                                                                                                                                                                                                                                                                                           0;
the administration, animal is protected from disease, where the animal comprises both CD4+ and CD8+ T cells. It is also useful for treating a disease (e.g. cancer, tumour, melanoma, thymoma, lymphoma, sarcoma, lung cancer, non-Hodgkin's lymphoma, leukaemia, Hodgkin's lymphoma, uterine cancer, cervical cancer, bladder cancer, kidney cancer, adenocarcinoma, breast cancer, prostate cancer, ovarian cancer and pancreatic cancer). The animal is further subjected to a cancer treatment including surgery, radiation, chemotherapy or gene therapy. The administration of (I), preferably dendritic cell is prior to, subsequent to or concurrent with, the cancer treatment. The present sequence is a tumour antigen derived epitope for inclusion in the composition of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antigen specific T-cell; detection; diagnosis; cancer specific T-cell; cancer; tumour; cervical cancer; prostate cancer; cellular immunity; immune therapy; cytostatic; immunostimulant; vaccine; antigenic peptide; human; human leukocyte antigen; HLA-A24 epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A detection method of antigen specific T-cells, comprises the use of plural antigenic peptides, useful in semi-quantitative determination of cancer specific T-cell frequencies and for monitoring cellular immunity.
                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                   Score 46; DB 5; Length 9;
Pred. No. 1.7e+06;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human CypB HLA-A24 epitope, SEQ ID NO:23.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABR84373 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-NOV-2000; 2000JP-00345094
                                                                                                                                                                                                                                                                                                      92.0%;
88.9%;
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                                                                                                                                                                                                                                                                                                                          Local Similarity 88.9
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                                                                                                                                                                                                                                                                                                                                                                                    1 KYHRVIKDF
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                                                                                                                                                                                                                                                               Sequence 9 AA;
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                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                   cytostatic; antiallergic; vaccine; allergic reaction suppressor; antigen;
allergy; epitope; cancer; immunoglobulin E antibody; desensitization;
cyclophilin B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Desensitizers or allergic reaction suppressors containing peptides originating from the same antigenic substances as reaction-inducing substances, useful for preventing or treating type I allergic diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                            0; Indels
               Length
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Pred. No. 1.7e+06;
1; Mismatches 0;
                              7e+06;
             92.0%; Score 46; DB
88.9%; Pred. No. 1.7e
iive 1; Mismatches
                                                                                                                                                                                                                                                                                Cyclophilin B protein amino acids 84-92.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 8; SEQ ID NO 1; 49pp; Japanese.
                                                                                                                                                                                      ADC17701 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1;
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88.9%;
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                                                                                                                                                                                                                                                    (first entry)
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                                               Conservative
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Query Match
Best Local Similarity
8, Conserve
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KFHRVIKDF
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Matches 8; Conser
                                                                           1 KYHRVIKDF
                                                                                                        KFHRVIKDF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ITOH/) ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9 AA;
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                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                    ADC17701;
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RESULT

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This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The peptides AAR96424-R96524 represent novel peptides derived from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hepatitis C virus; subtype; polymerase chain reaction; amplification; PCR; primer; probe; antibody; infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hepatitis C virus poly:nucleic acid unique to unidentified sub:type -used to develop probes and primers for new sub:types and vaccines to prevent and treat infection.
                                                                                                                                                                           Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                         Score 36; DB 3; Length 9; Pred. No. 1.7e+06; 0; Mismatches 1; Indels
                                                                                                                                                                                                                Claim 10; Page 59; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR96521 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis C virus type 9 peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 5; Page 68; 150pp; English.
                                                                                             SUMU ) SUMITOMO PHARM CO LTD
                                                                      98JP-00178449.
                                                                                                                                                                                                                                                                                                                        72.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95WO-EP004155
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95EP-00870076
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                                                                                                                                                                                                                                                                                                                                                  Conservative
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Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                        1 KYHRVIKD
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                                                                                                        ITOH K.
                                                                                                                                                                                                                                                                                                   Sequence 9 AA;
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WO9967288-A1
                                                                     25-JUN-1998;
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                                               24-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-OCT-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Maertens G,
                       29-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-MAY-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR96521;
                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                        (ITOH/)
                                                                                                                                Itoh K,
                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 11
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                                                                                                                                                                                                                                                                                                                                 Identifying a cell that differentiates into a metastatic cancer cell, useful for preventing metastatic cancer, comprises identifying a mutated fibronectin in the cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                       The invention comprises a method for identifying a cell that will differentiate into a metastatic cancer cell, the method involves identifying a mutated fibronectin in the cell. The method of the invention is useful for preventing metastatic cancer. The present acid sequence represents a Class I HLA-restricted widely expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; 0
                                                                                                      metastatic cancer cell differentiation; mutated fibronectin; metastatic cancer; class I HLA-restricted; widely antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 46; DB 8; Length 9;
Pred. No. 1.7e+06;
1; Mismatches 0; Indels
                                                                             Class I HLA-restricted widely expressed antigen #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label= Phe, Leu, Ile, Trp, Met
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /label= Phe, Tyr, Met, Trp
                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 84; 137pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human cyclophilin B peptide fragment #40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
         ADG89641 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY69960 standard; peptide; 9 AA
                                                                                                                                                                                                                                                             (BAYU ) BAYLOR COLLEGE MEDICINE
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88.9%;
                                                                                                                                                                                                                28-MAY-2003; 2003WO-US016736
                                                                                                                                                                                                                                       28-MAY-2002; 2002US-0383530P.
                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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KFHRVIKDF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 9 AA;
                                                                                                                                          Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                         11-MAR-2004
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                                  ADG89641;
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                                                                                                                                                                                                                                                                                     Wang R;
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Matches
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ADG89641
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Gaps

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novel hepatitis C virus subtypes 1d-f, 2e-i, 2k, 2l, 3g, 4k-m, 7a-c or types 9, 10 or 11 (see AATZ7937-TZ7989). The sequences corresp. to the 5' untranslated region (UR), the Core/El, NS4 or NS5B regions of the genome. This sequence is from the HCV subtype 9. The new HCV types were isolated from patients with chronic HCV from the Benelux countries, France, Cameroon and Vietnam, because of their aberrant reactivities. The RNA was extracted, cDNA synthesised and PCR amplified, cloned and genotyped. The 5'UR, Core/El and NS5B regions were sequenced either directly or partially and used to classify the new viruses into (sub)types based on comparison with known sequences. The nucleotide sequences can be used to synthesise probes and primers for the detection of HCV in a sample. The polypeptides can be used to detect anti-HCV antibodies, for HCV typing or to prevent HCV infections
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Sequence 10 AA;

. 0 62.0%; Score 31; DB 2; Length 10; 62.5%; Pred. No. 31; 1; Indels ed. No. 31; Mismatches Conservative KYHQVTQD 10 1 KYHRVIKD 8 Best Local Similarity Matches 5; Conserv Query Match g ð

AAB82292 standard; peptide; 11 AA.

(first entry) 09-JUL-2001 AAB82292;

3-Hydroxy-3-methylglutaryl-CoA reductase; HMG-CoA reductase; HMGR; phosphorylation; isoprenoid; sterol; transgenic plant; potato; pathogen resistance; disease resistance; crop protection.

Phosphorylation site in potato HMG-CoA reductase.

Solanum tuberosum

/note= "O-phosphorylated" Location/Qualifiers Key Modified-site

WO200131043-A1

03-MAY-2001

27-OCT-2000; 2000WO-GB004141.

99GB-00025453 27-OCT-1999;

(UYBR-) UNIV BRISTOL.

Beale MH; Hey SJ, Halford NG,

WPI; 2001-308654/32.

Modified 3-hydroxy-3-methylglutaryl-coenzyme A reductase gene, useful for generating transgenic plants having increased isoprenoid production, comprising a modified phosphorylation site.

Claim 16; Page 18; 23pp; English

The present sequence is that of a phosphorylated region of the 3-hydroxy-3-methylglutaryl-coenzyme A reductase (HMGR) protein of potato. The invention relates to modified HMGR genes useful for generating transgenic plants having increased isoprenoid (especially sterol) production. The modified genes encode modified HMGR proteins, in which a phosphorylation site has been rendered inactive by replacing a Ser, Thr or Tyr residue with Ala or other amino acid residue. This reduces transcriptional regulation. The HMGR genes may be in the form of mutant plant and 

regulation can also be avoided by using heterologous promoters. Increased seed starol content has been demonstrated in transgenic plants. The seed starol content has been demonstrated in transgenic plants. The invention is used to produce plants having increased insect and mite pest resistance, particularly against species of Homoptera, Diptera, Clepidoptera, Coleoptera, Hemiptera, Dictyoptera, Orthoptera, Coleoptera, Coleoptera, Nemenoptera, Nemenoptera, Nemenoptera, Nemenoptera, Coleoptera, Nemenoptera, Downy mildews, Colletotrichum, Cochliobolus, Tapesia, Magnaporthe, Stagnospora, Rhynchosporium, Septoria, Helminthosporium, and powdery mildews such as Blumeria and Erysiphe. The modified genes are particularly useful in maize, wheat, rice, barley, oilseed rape, beans, sunflower, cabbage, potato, spinach, broccoli, pea, cauliflower, tomato, forest trees, roses and tea. The invention may also be used to add nutritional benefit such as fat-soluble vitamins E and K, and sterols to a crop plant Transcriptional plant/non-plant or different plant chimeric genes. 

Sequence 11 AA;

Gaps

Gaps 0 62.0%; Score 31; DB 4; Length 11; 75.0%; Pred. No. 34; 1; Indels iive 1; Mismatches 1; Indels 6; Conservative Query Match Best Local Similarity Matches 6; Conserv

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1 KYHRVIKD 8

à g

KYNRSIKD 10

RESULT 13 AAY69959

AAY69959 standard; peptide; 9 AA.

AAY69959;

(first entry) 11-APR-2000 Human cyclophilin B peptide fragment #39

Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.

Homo sapiens

WO9967288-A1

29-DEC-1999.

99WO-JP003360. 24-JUN-1999;

25-JUN-1998;

(SUMU ) SUMITOMO PHARM CO LTD. (ITOH/) ITOH K.

Gomi S; rtoh K,

WPI; 2000-116932/10.

Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors. 

Claim 4; Page 61; 64pp; Japanese.

This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours

Sequence 9 AA;

Query Match

Length 9; DB 3; 58.0%; Score 29;

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Polyoma virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                  Gaps
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Pred. No. 83;
1; Mismatches 0; Indels
      Pred. No. 1.7e+06;
1; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HLA antigen; diagnosis; tumour; therapy
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83.3%;
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The invention relates to isolated Bacillus anthracis and small pox virus pebtides, also known as replikins, comprising 7 to 50 amino acids including at least one lysine residue located six to ten residues from a second lysine residue, at least one histidine residue, and at least 6% lysine residues. The peptides were identified using the 3-point.

Crecognition method and based on the replikin sequence isolated from glioblastoma cells malignin peptide which is of non-human origin. Also included are anti-replikin antibodies (or antibody cocktail), a method of stimulating the immune system of a subject to produce antibodies to Bacillus anthracis or more Bacillus anthracis or small pox comprising administering an effective amount of at least one or more Bacillus anthracis polypeptide (e.g. Anthrax Lethal Factor Protein pX01-07) comprising a replikin sequence, an antisense nucleic acid molecule complementary to a mRNA encoding a Small Pox Virus polypeptide (e.g. Surface Antigen S Precursor Protein comprising a replikin sequence. The produce antibodies that bind specifically to Anthrax or small pox provention comprising antibodies that can be used in the treatment and or prevention and or produce the peptides are useful for stimulating the treatment are useful for subject to produce antibodies that can be used in the treatment are useful for the preprides are usef
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolated Bacillus anthracis and small pox virus peptides (I), also known as replikins, useful for formulating vaccines for treating anthrax or small pox, comprise 7 to 50 amino acids.
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27-MAR-2001; 2001US-0278761P.
09-JUL-2001; 2001US-0303396P.
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5; Conservative
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(BOGO/) BOGOCH E
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Sequence 14, Appl
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                                                                                                                                                            October 21, 2004, 07:13:10 ; Search time 21.5 Seconds (without alignments) 27.761 Million cell updates/sec
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2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-08-973-335-11

US-08-467-976-31

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US-09-878-281A-264
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Maximum Match 100%
Listing first 45 summaries
                                                                                                            - protein search, using sw model
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#4.0   12   1   1   1   1   1   1   1   1	US-07-778-233B-29 US-08-290-641-29 US-08-290-641-29 US-08-548-540-29 US-08-742-243-6 US-08-742-243-9 US-08-742-243-9 US-08-742-243-9 US-08-772-243-9 US-08-877-605-93 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-605-94 US-08-877-978-37 US-08-877-978-37	ALIGNMENTS	S OF HEPATITIS C E AS PROPHYLACTIC S ASCII text outpu 075A 4155 6.9
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Query Match 62.0%; Score 31; DB 3; Length 10; Best Local Similarity 62.5%; Pred. No. 10; Matches 5; Conservative 2; Mismatches 1; Indels

JENGTH: 10 amino acid;
TYPE: amino acid;
TOPOLOGY: linear;
MOLECULE TYPE: peptide
US-08-836-075A-204

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TYPE: amino acid
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                                                                                                                                                                                                          COUNTRY: US
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APPLICANT: Jones, Michael H.

TITLE OF INVENTION: TRANSCRIPTIONAL REGULATOR
FILE REFERENCE: 06501-043001

CURRENT PILING DATE: 1999-10-15

PRIOR APPLICATION NUMBER: BCT/JP98/01782

PRIOR FILING DATE: 1998-04-17

PRIOR FILING DATE: 1997-04-18

NUMBER OF SEQ ID NOS: 28

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 22

LENGTH: 11
                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: JORES, Michael H.
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATOR
FILE REFERENCE: 06501-043001
CURRENT APPLICATION NUMBER: US/09/418,780A
CURRENT APPLICATION NUMBER: PCT/JP98/01782
PRIOR APPLICATION NUMBER: PCT/JP98/01782
PRIOR FILING DATE: 1998-04-17
PRIOR FILING DATE: 1997-04-18
NUMBER OF SEQ ID NOS: 28
NUMBER OF SEQ ID NOS: 28
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US-09-418-780A-14
; Sequence 14, Application US/09418780A
; Patent No. 6504009
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US-08-471-286-1
; Sequence 1, Application US/08471286
; Patent No. 5866688
; GENERAL INFORMATION:
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; ORGANISM: Drosophila melanogaster
US-09-418-780A-22
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Best Local Similarity 66.7
Matches 4; Conservative
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Best Local Similarity 66.7
Matches 4; Conservative
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ORGANISM: Homo sapiens
                                    KYHQVTQD 10
KYHRVIKD 8
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1 YHKIIK 6
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1 YHKIIK
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APPLICANT CHARLESKY, JOSEPH APPLICANT KIN, MEMORY T.
APPLICANT KIN, BOWARD T.
TITLE OF FUNDATION TO EXPONENTS
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MURRES OF SECURIORS
ANDESSERS RAINED CHARLESKY
MURRES OF SECURIORS
ANDESSERS RAINED CHARLESKY
MURRES TARGET MACHINE SECURIOR STORY
MURRES TARGET MACHINE SOUND STORY
MURRES APPLICATION NUMBER 10,956
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APPLICANT: Oppenheim, Frank G.
APPLICANT: Xu, Tao
APPLICANT: Spacciapoli, Peter
APPLICANT: Spacciapoli, Peter
APPLICANT: Roberts, F. D.
APPLICANT: Friden, Philip M.
TITLE OF INVENTION: Anti-Fungal D-Amino Acid Histatin-Based
TITLE OF INVENTION: Peptides
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
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    Indels
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ZIP: 02173

ZIP: 02173

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,563A
FILING DATE: 07-UUN-1996

CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/485,273
FILING DATE: 07-UUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: BROOK, David E.
REFERENCE/DOCKET NUMBER: PER95-02A2
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
TELECOMMUNICAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "At least one amino acid
must have a D configuration."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 50.0%; Score 25; DB 2; Le Best Local Similarity 100.0%; Pred. No. 1.7e+02; Matches 4; Conservative 0; Mismatches 0;
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;
  Mismatches
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; Sequence 31, Application US/08973559
; Patent No. 5912230
    4; Conservative
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OTHER INFORMATION:
OTHER INFORMATION:
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US-08-973-563A-31
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Sequence 8, Application PC/TUS9611495
GENERAL INFORMATION:
APPLICANT: UNIVERSITY OF MASSACHUSETTS MEDICAL CENTER
TITLE OF INVENTION: NUCLEAR LOCALIZED TRANSCRIPTION FACTOR
TITLE OF INVENTION: KINASES AND DIAGNOSTIC ASSAYS RELATED THERETO
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
CITY: Boston
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STATE: MA
COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/11495
FILING DATE: 03-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 25; DB 1; I
Pred. No. 1.5e+02;
1; Mismatches 1;
                    NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 04020/080001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Fasse, J. Peter
REGISTRATION NUMBER: 32,983
REFERENCE/DOCKET NUMBER: 04020/080WO1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
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APPLICATION NUMBER: US 08/503,062
FILING DATE: 10-JUL-1995
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                         TELEX: 200154
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-503-062-8
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Best Local Similarity
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Best Local Similarity
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1 YHDIIK 6
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Length 12;

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DB 3, Le.
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                                 ilarity 100.0%; Score 25; DB 10arity 100.0%; Pred. No. 1.7 Conservative 0; Mismatches
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CRGANISM: Homo sapiens
US-08-993-235-11
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Best Local Similarity
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Best Local Similarity
                                                     Best Local Similarity
Matches 4; Conserv
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US-10-037-311B-16
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US-09-226-666-7
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LENGTH: 12
                                     Query Match
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; Faquence 11, Application US/08993235
; Patent No. 6084064
; GENERAL INFORMATION:
; APPLICANT: OPPENHEIM, FRANK G.
; APPLICANT: ROBERTS, F. DONALD
; APPLICANT: XU, TAO
; APPLICANT: XU, TAO
; APPLICANT: XU, TAO
; TILLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES
; FILE REFERENCE: 50032/002001
; FILE REFERENCE: 1997-12-18
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 11
; LENGTH: 12
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APPLICANT: XU, TAO
APPLICANT: XU, TAO
APPLICANT: ROBERTS, F. D.
APPLICANT: SPACCIAPOLI, PETER
APPLICANT: FRIDEN, PHILIP M.
TITLE OF INVENTION: Anti-Fungal and Anti-Bacterial
TITLE OF INVENTION: Histatin-Based Peptides
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
STATE: MA
                                                                                                                                                                                                                                                                                                                 COMPUTER EADABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,559
FILING DATE: 07-JUN-1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/481,888
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-01A2
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 25; DB 2; Lot Pred. No. 1.7e+02;
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid STRANDEDNESS:
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Best Local Similarity
Matches 4; Conserv
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Fatent No. 6531573
GENERAL INFORMATION:
APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: SPACICIAPOLI, PETER
TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES
FILE REFERENCE: 50032/002001
CURRENT FILING DATE: 1997-12-18
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 3.0
; Sequence 7. Application US/09226666A; Sequence 7. Application US/09226666A; Patent No. 6528488; GENERAL INFORMATION:
; APPLICANT: Spacciapoli, Peter; APPLICANT: Rothstein, David M.; TITLE OF INVENTION: METHOD FOR TREATING CYSTIC FIBROSIS; FILE REFERENCE: 50032/007001; CURRENT APPLICATION NUMBER: US/09/226,666A; CURRENT FILING DATE: 1999-01-08; NUMBER OF SEQ ID NOS: 18; SOFTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 50.0%; Score 25; DB 4; Length 12; 100.0%; Pred. No. 1.7e+02; cive 0; Mismatches 0; Indels
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us-09-720-469a-39.rai

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SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
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STRANDEDNESS: single
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                                       1 KYHRVIKD 8
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Best Local Similarity
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APPLICANT: HEY, TIMOTHY D
APPLICANT: HEY, 
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COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/378,761A
FILING DATE: 26-JAN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
        Sequence, Sequence, Patent No. 6747190

Patent No. 6747190

GENERAL INFORMATION:
APPLICANT MICHIGAN STATE UNIVERSITY

ITLE OF INVENTION XYLOGLUCAN FUCOSYLTRANSFERASES
FILE REFERENCE: MS00-00102;
CURRENT APPLICATION NUMBER: US/10/037,3118

CURRENT FILING DATE: 2001-11-09

PRIOR APPLICATION NUMBER: US60/117,555

PRIOR FILING DATE: 1999-01-28

NUMBER OF SEQ ID NOS: 17

SOFTWARE: PatentIn version 3.1

SEQ ID NO 16

LENGTH: 12
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REGISTRATION NUMBER: 33651
REFERENCE/DOCKET NUMBER: 38272B
TELECOMMUNICATION INFORMATION:
TELEPHONE: (317) 337-4846
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-378-761A-68
; Sequence 68, Application US/08378761A
; Patent No. 5635384
; GENERAL INFORMATION:
; APPLICANT: WALSH, TERENCE A
; APPLICANT: HEY, TIMOTHY D
Sequence 16, Application US/10037311B
Patent No. 6747190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               50.0%; Scor.
66.7%; Pred
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STREET: 9330 ZIONSVILLE ROAD
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CITY: INDIANAPOLIS
STATE: IN
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Best Local Similarity
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US-08-485-286-68

Sequence 68, Application US/08485286

Patent No. 5646026

Patent No.
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; APPLICANT: Spacciapoli, Peter; APPLICANT: Rothstein, David M.; APPLICANT: Friden, Phillip M.; TITLE OF INVENTION: METHOD FOR TREATING CYSTIC FIBROSIS; FILE REFERENCE: 50032/007001; FILE REFERENCE: 50032/007001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/485,286 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 26-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: BORUCKI, ANDREA T
REGISTRATION NUMBER: 33651
REFERENCE/DOCKET NUMBER: 38272B
TELECOMMUNICATION INFORMATION:
TELEPHONE: (317) 337-4846
INFORMATION FOR SEQ ID NO: 68:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 09/378761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 46268
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CURRENT FILING DATE: 1999-01-08

CURRENT FILING DATE: 1999-01-08

NUMBER OF SEQ ID NOS: 18

SOFTWARE: FastSEQ for Windows Version 4.0

EDST ID NO 17

LENGTH: 12

TYPE: PRT

ORGANISM: Artificial Sequence
FRATURE:

COTHER INFORMATION: Derived from Homo sapiens

OCHER INFORMATION: Derived from Homo sapiens

SOURCE STATE INFORMATION: Derived from Homo sapiens

OUGLY MATCH

Best Local Similarity 44.4%; Score 23; DB 4; Length 12;

Best Local Similarity 44.4%; Pred. No. 3.9e+02;

MATCHES 4; Conservative 2; Mismatches 3; Indels 0;

OY 1 KYHRVIKDF 9

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Search completed: October 21, 2004, 07:29:30 Job time: 22.5 secs

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Sequence 1, Appli
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Sequence 1, Appli
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                                                                                                                                                                             October 21, 2004, 07:15:25 ; Search time 61.75 Seconds (without alignments)
47.187 Million cell updates/sec
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1: cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
2: cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
4: cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
5: cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
6: cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
7: cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
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9: cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
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GenCore version 5.1.6
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US-09-984-057-61

US-09-984-057-61

US-10-105-232-61

US-10-105-232-62

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# ALIGNMENTS

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                                                                      PEPLICANT: Wang, Rong-fu
TITLE OF INVENTION: Mutant Fibronectin and Tumor Metastasis
FILE REFERENCE: HO-P02484US1
CURRENT APPLICATION NUMBER: US/10/447,161
CURRENT FILING DATE: 2003-05-28
PRIOR APPLICATION NUMBER: 60/383,530
PRIOR FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 148
SOFTWARE: Patentin version 3.1
SEQ ID NO 84
                                                                                                                                                                                                                                                                                                                                                                                                           92.0%; Score 46; DB 15; Length 9; 88.9%; Pred. No. 1.2e+06;
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                  Sequence 84, Application US/1047161
Publication No. US20040023314A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence 1, Application US/10788016; Publication No. US20040141992A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
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Pred. No. 64;
2; Mismatches 0; Indels
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               DB 9; Length 10; 24;
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j Bublication No. US20030083470A1

general information:
TITLE OF INVENTION:
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATOR
FILE REFERENCE: 36501-043001
CURRENT APPLICATION NUMBER: US/10/293,822
CURRENT FILING DATE: 2002-11-12

PRIOR PRICATION NUMBER: US/09/418,780

PRIOR FILING DATE: 1999-10-15

PRIOR FILING DATE: 1999-10-15

PRIOR FILING DATE: 1999-04-17

PRIOR FILING DATE: 1997-04-18

NUMBER OF SEQ ID NOS: 28

SOFTWARE: FastSEQ for Windows Version 4.0

TENNOME 11
                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:

APPLICANT: Jones, Michael H.

TITLE OF INVENTION: TRANSCRIPTIONAL REGULATOR
FILE REFERENCE: 06501-043001

CURRENT APPLICATION NUMBER: US/10/293,822

CURRENT FILING DATE: 2002-11-12

PRIOR FILING DATE: 1999-10-15

PRIOR FILING DATE: 1999-04-17

PRIOR FILING DATE: 1998-04-17

PRIOR FILING DATE: 1998-04-17

PRIOR FILING DATE: 1998-04-17

PRIOR FILING DATE: 1997-04-18

NUMBER OF SEQ ID NOS: 28

SOFTWARE: FastSEQ for Windows Version 4.0

LENGTH: 11

LENGTH: 11
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                                                           2; Mismatches
                 62.0%; Score 31; 62.5%; Pred. No. 2
                                                                                                                                                                                                                                                        ; Sequence 14, Application US/10293822; Publication No. US20030083470A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Drosophila melanogaster US-10-293-822-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        58.0%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 66.7
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Best Local Similarity 66.7
Matches 4; Conservative
                 Query Match
Best Local Similarity 62.5
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
                                                                                                                                                KYHQVTQD 10
                                                                                                      1 KYHRVIKD 8
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1 YHKIIK 6
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                                                                                                         δ
                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Peptide consisting of 9 amino acid residues from the 84th residue of OTHER INFORMATION: to the 92nd residue of cyclophillin B
US-10-788-016-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STUYVER, LIEVEN
TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES
AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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ZIP: 77210-4433

COUNTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 6.0 / ASCII text output
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/851,138
FILING DATE: 09-May-2001
PRIOR APPLICATION NUMBER: 08/836,075
FILING DATE: <UNKNOWN>
APPLICATION NUMBER: EP 94870166.9
FILING DATE: 21 Oct 1994
APPLICATION NUMBER: EP 95870076.7
FILING DATE: 22 Jun 1995
                                                                                                                                                                                                                                                                                                                                                                                                          Score 46; DB 16; Length 9; Pred. No. 1.2e+06; 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 207
CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE & DURKEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: KAMMERER, PATRICIA A.
REGISTRATION NUMBER: 29,775
REFERENCE/DOCKET NUMBER: INNS:004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
TOPOLOGY: linear
;
MOLECULE TYPE: peptide
;
SEQUENCE DESCRIPTION: SEQ ID NO: 204:
US-09-851-138-204
                                                           CURRENT FILING DATE: 2004-02-26
PRIOR APPLICATION NUMBER: PCT/JP02/08641
PRIOR FILING DATE: 2002-08-28
PRIOR APPLICATION NUMBER: JP P2001-260046
PRIOR FILING DATE: 2001-08-29
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.2
                   FILE REFERENCE: 3190-049
CURRENT APPLICATION NUMBER: US/10/788,016
CURRENT FILING DATE: 2004-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 204, Application US/09851138 Publication No. US20020183508A1 GENERAL INFORMATION:
TITLE OF INVENTION: Desensitizers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: MAERTENS, GEERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: P.O. BOX 4433
                                                                                                                                                                                                                                                                                                                                                                                                              92.0%;
88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 88.9
----hes 8; Conservative
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KFHRVIKDF 9
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                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Artificial
FEATURE:
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US-09-851-138-204
                                                                                                                                                                                                                 SEQ ID NO 1
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APPLICANT: BOGOCH, SANDEL
APPLICANT: BOGOCH, ELENORE S.
TITLE OF INVENTION: REPLIKINS AND METHODS OF IDENTIFYING
TITLE OF INVENTION: REPLIKIN-CONTAINING SEQUENCES
FILE REFERENCE: 09425-46902
CURRENT APPLICATION NUMBER: US/09/984,057
CURRENT FILING DATE: 2001-10-26
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR FILING DATE: 2001-07-09
PRIOR FILING DATE: 2001-03-27
PRIOR FILING DATE: 1998-09-04
PRIOR FILING DATE: 1998-09-04
PRIOR FILING DATE: 2001-03-27
PRIOR FILING DATE: 2001-03-27
PRIOR FILING DATE: 1994-02-17
NUMBER OF SEQ ID NOS: 90
SOFTWARE: PATENTIN 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    56.0%; Score 28; DB 9; 162.5%; Pred. No. 1.2e+06;
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Patent No. US20020151677A1
GENERAL INFORMATION:
APPLICANT: BOGOCH, SAMUEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 62.5
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TYPE: PRT
; ORGANISM: Polyoma virus
US-09-984-057-61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Polyoma virus
                         1 KYHRVIKD 8
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1 KQHRELKD 8
                                                                                  1 KOHRELKD
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US-09-984-057-62
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Sequence 62, Application US/09984056

Sequence 62, Application US/09984056

Patent No. US20020120106A1

GENERAL INFORMATION:

APPLICANT: BOGOCH, SAWUEL

APPLICANT: BOGOCH, ELENORE S.

TITLE OF INVENTION: ANTHRAX AND SMALL POX REPLIKINS AND METHODS OF USE

FILE REFERENCE: 09425-46903

CURRENT APPLICATION NUMBER: US/09/984,056

CURRENT FILING DATE: 2001-026

PRIOR FILING DATE: 2001-07-09

PRIOR FILING DATE: 2001-03-27

PRIOR FILING DATE: 1998-09-04

PRIOR FILING DATE: 1998-09-04

PRIOR FILING DATE: 1994-02-17

PRIOR FILING DATE: 1994-02-17

NUMBER OF SEQ ID NOS: 103

SEQ ID NO 62

LENGTH: BALCHT OF THE SECOND NOS: 103

LENGTH: BALCHT OF SEQ ID NOS: 103
                                                                                                                                                                                              GENERAL INFORMATION:

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: BOGOCH, SAMUEL

APPLICANT: BOGOCH, ELENORE S.

TITLE OF INVENTION: ANTHRAX AND SMALL POX REPLIKINS AND METHODS OF USE

FILE REFERENCE: 09425-46903

CURRENT APPLICATION NUMBER: 12001-10-26

PRIOR APPLICATION NUMBER: 60/303,396

PRIOR APPLICATION NUMBER: 60/278,761

PRIOR APPLICATION NUMBER: 09/146,755

PRIOR FILING DATE: 2001-03-27

PRIOR FILING DATE: 1998-09-04

PRIOR FILING DATE: 1998-09-04

PRIOR FILING DATE: 1994-02-17

PRIOR FILING DATE: 2001-03-27

PRIOR FILING DATE: 1994-02-17

NUMBER OF SEQ ID NOS: 103

SOFTWARE: PATENTIN 1.144
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                                                                                                                                               Sequence 61, Application US/09984056 Patent No. US20020120106A1
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Best Local Similarity 62.5
Matches 5; Conservative
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; ORGANISM: Polyoma virus
US-09-984-056-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Polyoma virus
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Best Local Similarity
Matches 5; Conserv
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                                                                                  RESULT 6
US-09-984-056-61
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Indels Length 9;

1; Mismatches

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Sequence 61. Application US/09984057

Sequence 61. Application US/09984057

Patent No. US20020151677A1

GENERAL INFORMATION:

APPLICANT: BOGOCH, SAMUEL

APPLICANT: BOGOCH, ELENORE S.

TITLE OF INVENTION: REPLIKINS AND METHODS OF IDENTIFYING

TITLE OF INVENTION: REPLIKIN-CONTAINING SEQUENCES

FILE REFERENCE: 09425-46902

CURRENT APPLICATION NUMBER: US/09/984,057

CURRENT FILING DATE: 2001-0-26

PRIOR FILING DATE: 2001-07-09

PRIOR FILING DATE: 2001-03-27

PRIOR FILING DATE: 1998-09-04

PRIOR FILING DATE: 1998-09-04

PRIOR FILING DATE: 1994-02-17

NUMBER OF SEQ ID NOS: 90

SOFTWARE: Patentin 2.1

SEQ ID NO 61
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us-09-720-469a-39.rapb

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SOFTWARE: Patentin 2.1 SEQ ID NO 62
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US-10-189-437-62
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US-10-105-232-61

Sequence 61, Application US/10105232

Publication No. US20030180328A1

GENERAL INFORMATION:
APPLICANT: BOGOCH, SAMUEL

APPLICANT: BOGOCH, ELENDER S.

TITLE OF INVENTION: AND IN INFLUENZA EPIDEMICS

FILE REPRESENCE: 09425-46904

CURRENT PILING DATE: 2002-03-26

PRIOR PRIOR DATE: 2001-07-09

PRIOR PPLICATION NUMBER: 60/303,396

PRIOR PLING DATE: 2001-03-27

PRIOR PLING DATE: 2001-03-27

PRIOR APPLICATION NUMBER: 09/146,755

PRIOR APPLICATION NUMBER: 09/146,755

PRIOR PPLING DATE: 2001-03-27

PRIOR APPLICATION NUMBER: 08/198,139

PRIOR PLING DATE: 2001-03-27

PRIOR APPLICATION NUMBER: 08/198,139

PRIOR PRIOR FILING DATE: 2001-03-27

PRIOR PRIUMG DATE: 1994-02-17

NUMBER OF SEQ ID NOS: 535

SOFTWARE: PATENTIN 2.1

SRQ ID NOS: 535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 62, Application US/10105232

Publication No. US20030180328A1

GENERAL INFORMATION:

APPLICANT: BOGOCH, SAMUEL

APPLICANT: BOGOCH, ELENORE S.

TITLE OF INVENTION: REPLIKIN PEPTIDES IN RAPID REPLICATION OF GLIOMA CELLS

TITLE OF INVENTION: AND IN INFLUENZA EPIDEMICS

FILE REFERENCE: 09425-46904

CURRENT APPLICATION NUMBER: 00/303,396

PRIOR APPLICATION NUMBER: 60/303,396

PRIOR FILING DATE: 2001-07-09

PRIOR APPLICATION NUMBER: 09/146,755

PRIOR FILING DATE: 1998-09-04

NUMBER OF SEQ ID NOS: 535
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                                              56.0%; Score 28; DB 9; Length 9; 62.5%; Pred. No. 1.2e+06; iive 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 62.5
Matches 5; Conservative
                                                                                                 Conservative
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; ORGANISM: Polyoma virus
US-10-105-232-61
                                                                                                                                            1 KYHRVIKD 8
                                                                                                                                                                                          KOHRELKD 8
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                                              Query Match
Best Local Similarity
Matches 5; Conserv
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US-10-105-232-62
 US-09-984-057-62
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Search completed: October 21, 2004, 07:33:43 Job time : 62.75 secs
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Sequence 76, Application US/10601953
Sequence 76, Application US/10601953
Sequence 100 No. US20040077540A1
GENERAL INFORMATION:
APPLICANT: Quay, Steven C.
TITLE OF INVENTION: Compositions And Methods For Modulating Physiology Of Epithelial
TITLE OF INVENTION: Therapeutic Compounds
FILLE REFERENCE: 02-03US
CURRENT APPLICATION NUMBER: US/10/601,953
CURRENT APPLICATION NUMBER: 60/392,512
PRIOR FILING DATE: 2002-06-28
NUMBER OF SEQ ID NOS: 900
SOFTWARE: Patentin version 3.2
SEQ ID NO 76
LENGTH: 12
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; Sequence 139, Application US/09834765
; Patent No. US20020055478A1
; GENERAL INFORMATION:
; APPLICANT: Mary Faris
; APPLICANT: Arthur B. Raitano
; APPLICANT: Steve Chappell Mitchell
; APPLICANT: Daniel E.H. Afar
APPLICANT: Aya Jakobovits
; TITLE OF INVENTION: GTP-BINDING PROTEIN USEFUL IN TREATMENT
; TITLE OF INVENTION: GTP-BINDING PROTEIN USEFUL
; TITLE OF INVENTION: AND DETECTION OF CANCER
; TITLE OF INVENTION: AND DETECTION OF CANCER
; CURRENT APPLICATION NUMBER: US/09/834,765
; CURRENT FILING DATE: 2000-04-12
; PRIOR FILING DATE: 2000-04-12
; NUMBER OF SEQ ID NOS: 770
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 139
; LENGTH: 9
                                                                                                                                                                                                      ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 56.0%; Score 28; DB 15; Length 12; Best Local Similarity 57.1%; Pred. No. 1.1e+02; Matches 4; Conservative 2; Mismatches 1; Indels
                                                                                                                                                Query Match 56.0%; Score 28; DB 14; Length 9; Best Local Similarity 62.5%; Pred. No. 1.2e+06; Matches 5; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Artificial Sequence
; SEQ ID NO 62
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Polyoma virus
US-10-189-437-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
CRGANISM: Homo sapiens
US-09-834-765-139
                                                                                                                                                                                                                                                        1 KYHRVIKD 8
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1 KQHRELKD 8
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1 HNIIQDF 7
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US-10-601-953-76
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54.0%; Score 27; DB 9; Length 9; 62.5%; Pred. No. 1.2e+06; iive 1; Mismatches 2; Indels
                               5; Conservative
                                                           2 YHRVIKDF 9
   Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                      1 YEHVIKNF
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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using sw model - protein search, OM protein

Run on:

October 21, 2004, 07:12:54; Search time 15.75 Seconds (without alignments) 54.981 Million cell updates/sec

US-09-720-469A-39 50 1 KYHRVIKDF 9

score: Sequence: Title: Perfect

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

283416 segs, 96216763 residues Searched: 1582 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 12

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

PIR_79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

RESULT 2

SUMMARIES	Description	Ig heavy chain V r	telomeric and tetr	Rhesus blood group	'O	hypothetical 1.3K	cytochrome-c oxida	glycine reductase	T-cell receptor al		ž	virG protein - Agr	peptide-N4- (N-acet	cycloinulooligosac	nrc		٠,	unidentified 5.4/3			Na+-transporting A	hutU protein - Kle		calliFMRFamide 10	bone gla protein -	serum heterodimer,	proteasome endopep	unidentified 6.0/1	T antiden variant	· · · · · · · · · · · · · · · · · · ·
SUMIN	ID	536849	PC4377	154193	PC4374	JQ0943	\$65381	B39308	S23373	G64003	NYPG7	S13224	A59272	PC2330	152708	S10624	PT0228	PQ0730	B44510	PT0246	866296	C36730	878420	A44787	149406	B56899	G42762	PQ0733	PH1376	
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o ^k	Query Match		38.0	'n	4	34.0	32.0	ď	ζ,	ς.	0	0	30.0	0	30.0	0	30.0	0	28.0	œ	æ	28.0	28.0	28.0	80	28.0	8	28.0	28.0	•
	Score	20	19	18	17	17	16	16	16	16	15	15	15	15	15	15	15	15	14	14	14	14	14	14	14	14	14	14	14	1
	Result No.	1 1 1 1 1 1	7	e	4	Ŋ	9	7	8	σ	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	. 56	27	28	)

proteasome chain 1 S34447 S34447 binR protean 1 S02617 S02617 S02617 S02617 S03165 S03165 Foredoxin a2 - Ja Hypothetical prote For Foredoxin a2 - Ja Hypothetical prote For Foredoxin a2 - Ja Hypothetical prote For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD PPT0315 For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD PPT0315 For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD PPT0315 For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD PPT0315 For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD For Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypothetical prote Igheavy chain CRD Foredoxin a2 - Ja Hypoth	ALIGNMENTS  use)  ision 01-Dec-1995 #text_change 05-Nov-1999  f, July 1992  the primary immune response to (4-hydroxy-3-nitropheny)  NID:g50070; PIDN:CAA47794.1; PID:e51590; PID:g1333861  oglobulin  Score 20; DB 2; Length 10;  Pred. No. 9.1e+02;  3; Mismatches 3; Indels 0; Gaps 0;	
\$3940 \$3340 \$302 \$565 \$656 \$740 \$740 \$741 \$741 \$741 \$741 \$741 \$741 \$741 \$741	- mouse house) (house mouse) uence_revision ta Library, Jul tudies on the B 24 acid acid .:X67382; NID:gl er; immunoglobi 40.0%; Scorr 33.3%; Pred ative 3; M	
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H H H A A A A A A A A A A A A A A A A A	- mou (hous nence ta Li tudie 24 .:X673 her; i er; i er; i	9 10
0 14 28.0 1 14 28.0 2 13 26.0 4 13 26.0 4 13 26.0 7 13 26.0 7 13 26.0 9 13 26.0 1 13 26.0 1 13 26.0 1 13 26.0 1 13 26.0 1 13 26.0 2 13 26.0 3 13 26.0	chain V region " Mus musculus 6-Jan-1994 #seq on: \$36849 J.; Kelsoe, G. I to the EMBL Da ption: In situ s ption: In situ s to re number: \$250 on: \$36849 on: \$36849 er type: nucleic ss: 1-10 <jac> references: EMBL 8s: heterotetram Aatch</jac>	1 KYHRVIKDF :   : :  2 RYRTVVVNF
6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	RESULT 1 836849 19 heavy C;Species C;Date: C;Accessi R;Jacob, submitted A;Descrip A;Referer A;Referer A;Residue A;Cross-1 C;Reyword Query N Best LG	oy G

telomeric and tetraplex DNA binding protein qTBP42 VIII - rat (fragment)

C; Species: Rattus norvegicus (Norway rat)
C; Date: 28-Oct-1997 #sequence_revision 28-Oct-1997 #text_change 09-Jul-2004
C; Date: 28-Oct-1997 #sequence_revision 28-Oct-1997 #text_change 09-Jul-2004
C; Date: 28-Oct-1997 #sequence_revision 28-Oct-1997 #text_change 09-Jul-2004
R; Sarig, G:, Weisman-Shomer, P.; Fry, M.
R; Sarig, G:, Weisman-Shomer, P.; Fry, M.
Biochem. Biophys. Res. Commun. 237, 617-623, 1997
A; Title: Telomeric and tetraplex DNA binding properties of qTBP42: A homologue of the CAn A; Reference number: PC4371; MUID:97445086; PMID:9299414
A; Reference number: PC4371
A; Molecule type: protein
A; Residues: 1-12 <SAR>
A; Residues: 1-12 <SAR>
C; Comment: This protein binds either strand of the telomeric DNA as well as unimolecular ; 0 Gaps 0; Score 19; DB 2; Length 12; Pred. No. 1.7e+03; 1; Indels h 38.0%; Similarity 60.0%; 3; Conservative 1 Query Match Best Local Similarity Matches 3; Conserv 1 KYHRV 5 ð

|:| | 3 KFHTV 7 g

RESULT 3

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C;Accession: S65381
R;Schaegger, H.; Noack, H.; Halangk, W.; Brandt, U.; von Jagow, G.
Eur. J. Biochem. 230, 235-241, 1995
A;Title: Cytochrome-c oxidase in developing rat heart. Enzymic properties and amino-termi
A;Reference number: S65372; MUID:95324529; PMID:7601105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Glycine reductase (EC 1.4.99.-) sulfhydryl protein C, beta chain - Clostridium sticklandi
C;Species: Clostridium sticklandii
C;Date: 19-Jun-1992 #sequence_revision 19-Jun-1992 #text_change 09-Jul-2004
C;Accession: B39308
R;Stadtman, T.C.; Davis, J.N.
J. Biol. Chem. 266, 22147-22153, 1991
A;Title: Glycine reductase protein C. Properties and characterization of its role in the A;Reference number: A39308, MUID:92042141; PMID:1939235
A;Reference number: A39308
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-10 <STA>
A;Cross-references: UNIPROT:Q9EV92
C;Function:
A;Description: glycine reductase complex catalyzes the reductive deamination of glycine t
C;Keywords: ATP; oxidoreductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              T-cell receptor alpha chain J region - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999
C;Accession: $23373
R;Pluschke, G; Ricken, G; Taube, H; Kroninger, S; Melchers, I; Peter, H.H.; Eichmanr Eur. J. Immunol. 21, 2749-2754, 1991
A;Title: Biased T cell receptor V(alpha) region repertoire in the synovial fluid of rheun A;Reference number: $23364; MUID:92037820; PMID:1657615
A;Accession: $23373
A;Accession: $23373
A;Kestus: preliminary; translation not shown
A;Residues: 1-11 <PLU>
                                                                                                                                                                                                                                                                                                                 C;Species: Rattus norvegicus (Norway rat)
C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
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66.7%;
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Best Local Similarity 50.0
Matches 3; Conservative
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A;Molecule type: protein
A;Residues: 1-8 <SCH>
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Matches 2; Conserv
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4 NKVIAD 9
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RYH 10
KYH 3
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S23373
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JQ0943
hypothetical 1.3K protein - infectious bursal disease virus (strain 52/70)
C;Species: infectious bursal disease virus
C;Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change 09-Jul-2004
C;Accession: JQ0943
R;Bayliss, C.D.; Spies, U.; Shaw, K.; Peters, R.W.; Papageorgiou, A.; Mueller, H.; Bours J. Gen. Virol. 71, 1303-1312, 1990
A;Title: A comparison of the sequences of segment A of four infectious bursal disease vi A;Reference number: JQ0941; MJD:90278420; PMID:2161902
A;Accession: JQ0943
A;Accession: JQ0943
A;Molecule type: Genomic RNA
A;Molecule type: Genomic RNA
A;Molecule type: Genomic RNA
A;Residues: 1-10 <BAY>
A;Cross-references: UNIPROT:Q82625; GB:D00869; NID:g221038; PIDN:BAA00743.1; PID:d100120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CiSpecies: Rattus norvegicus (Norway rat)
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CiSpecies: Rattus norvegicus (Norway rat)
CiSpecies: Rattus norvegicus (Norway rat)
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CiSpecies: Rattus norvegicus (Norway rat)
CiSpecies: Rattus norvegicus (Norway rat)
CiSpecies: CiBecies: Rattus norvegicus (Norway rat)
CiSpecies: Commun. 28.7, M.
Biochem. Biophys. Res. Commun. 237, 617-623, 1997
Biochem. Biophys. Res. Commun. 237, 617-623, 1997
A;Title: Telomeric and tetraplex DNA binding properties of qTBP42: A homologue of the CAA; Milecuse number: PC4371; MUID:97445086; PMID:9299414
A;Reference number: PC4371; MUID:97445086; PMID:9299414
A;Reference number: PC4371; Muid: 
         Rhesus blood group CcEe protein - human (fragment)
C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 06-Sep-1996 #sequence_revision 06-Sep-1996 #text_change 09-Jul-2004
C; Accession: I54193
R; Cherif-Zahar, B.; Le Van Kim, C.; Rouillac, C.; Raynal, V.; Cartron, J.P.; Colin, Y. Genomics 19, 68-74, 1994
A; Title: Organization of the gene (RHCE) encoding the human blood group RhCcEe antigens A; Reference number: I54193; MUID:94245182; PMID:8188244
A; Accession: I54193
A; Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                     A;Molecule type: DNA
A;Residues: 1-11 <RES>
A;Cross-references: UNIPROT:Q9UPC8; UNIPROT:Q99906; UNIPROT:Q9UK19; UNIPROT:Q9UFNO; UNIPROT:Q9UPNO; UNIPROT:Q9UFNO; UNIPROT:Q9UEC7; GB:S70456; NID:G546795; PIDN:AAD14061.1; PID:G4261761
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 36.0%; Score 18; DB 2; Length 11; 42.9%; Pred. No. 2.4e+03; Live 2; Mismatches 2; Indels
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34.0%; Score 17; DB 2; Langth 10;

Best Local Similarity 66.7%; Pred. No. 3.4e+03;

Matches 2; Conservative 1; Mismatches 0; indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Genetics:
A;Gene: GDB:RHCE
A;Cross-references: GDB:229957; OMIM:111700
A;Map position: 1p36.2-1p34
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A;Accession: PC2330
A;Molecule type: protein
A;Residues: 1-11 <KUS>
A;Cross-references: UNIPROT:Q7M0L3
C;Comment: This enzyme hydrolyzes beta-(2-1) glycosidic linkages and acts in intermolecul
C;Keywords: glycosyltransferase; hexosyltransferase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cispecies: Bacillus circulans
CiDate: 21-Mar-1995 #sequence_revision 26-May-1995 #text_change 09-Jul-2004
CiAccession: PC2330
RiKushibe, S.; Mitsui, K.; Yamagishi, M.; Yamada, K.; Morimoto, Y.
Biosci. Biotechnol. Biochem. 59, 31-34, 1995
A;Title: Purification and characterization of cycloinulooligosaccharide fructanotransferals, Reference number: PC2330; MUID:95201377; PMID:7765973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cycloinulooligosaccharide fructanotransferase (EC 2.4.-.-) - Bacillus circulans (MCI-2554
J. Mol. Biol. 215, 537-547, 1990
A;Title: Binding of the regulatory protein VirG to the phased signal sequences upstream
A;Reference number: S13224; MUID:91039316; PMID:2231718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide-N4-(N-acetyl-beta-glucosaminyl)asparagine amidase (EC 3.5.1.52) A, large chain
N;Alternate names: peptide N-glycosidase
C;Species: Prunus dulcis var. sativa (sweet almond)
C;Date: 19-May-2000 #sequence_revision 19-May-2000 #text_change 09-Jul-2004
C;Accession: A59272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Altmann, F.; Paschinger, K.; Dalik, T.; Vorauer, K.

Bur. J. Biochem. 252, 118-123, 1998
A;Title: Characterisation of peptide-N4-(N-acetyl-beta-glucosaminyl) asparagine amidase.
A;Reference number: A59272; MUID:98181894; PMID:9523720
A;Accession: A59272
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-10 <ALT>
A;Cross-references: UNIPROT:P81899
C;Keywords: hydrolase
                                                                                           A,Accession: S13224
A,Status: preliminary
A,Molecule type: protein
A,Residues: 1-10 <TAM>
A,Cross-references: UNIPROT:Q7M0P7
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C;Species: Sus scrofa domestica (domestic pig)
C;Species: Sus scrofa domestica (domestic pig)
C;Accession: A01417
R;Chang, R.C.C.; Huang, W.Y.; Arimura, A.; Redding, T.W.; Coy, D.H.; Saffran, M.; Kong, Horm. Metab. Res. 13, 228-232, 1981
A;Title: Isolation, structure and synthesis of a heptapeptide with in vitro ACTH-releasi
A;Reference number: A01417; MUID:81213980; PMID:6263778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C; Accession: C64003
K; Fleischmann, R.D.; Adams, M.D.; White, O.; Clayton, R.A.; Kirkness, E.F.; Kerlavage, F; Fleischmann, R.D.; Shirley, R.; Liu, L.I.; Glodek, A.; Kelley, J.M.; Weidman, J.D.; Scott, J.D.; Scott, J.D.; Fritchman, J.L.; Fuhrmann, J.L.; Geoghagen, N.S.M.; Science 269, 496-512, 1995
A; Authors: Gnehm, C.L.; McDonald, L.A.; Small, K.V.; Fraser, C.M.; Smith, H.O.; Venter, A; Title: Whole-genome random sequencing and assembly of Haemophilus influenzae Rd.
A; Reference number: A64000; MuID:95350630; PMID:7542800
A; Reterence nucleic acid sequence not shown; translation not shown
A; Status: nucleic acid sequence not shown; translation not shown
A; Molecule type: DNA
A; Residues: 1-12 < TIGR>
A; Cross-references: GB:U32704; GB:L42023; NID:g1573143; PID:g1573155; TIGR:HI0195
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virG protein - Agrobacterium sp. (fragment)
C;Species: Agrobacterium sp.
C;Date: 19-Mar-1997 #sequence_revision 01-Aug-1997 #text_change 09-Jul-2004
C;Accession: $13224
R;Tamamoto, $..; Aoyama, T.; Takanami, M.; Oka, A.
                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein H10195 - Haemophilus influenzae (strain Rd KW20)
C;Species: Haemophilus influenzae
C;Date: 18-Aug-1995 #sequence_revision 18-Aug-1995 #text_change 30-Jun-1998
C;Accession: G64003
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                                                                                         Score 16; DB 2; Length 11;
Pred. No. 5.9e+03;
1; Mismatches 1; Indels
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A; Molecule type: protein
A; Residues: 1-7 < CHA>
A; Cross-references: UNIPROT: P01153
C; Superfamily: hypothalamic heptapeptide
C; Keywords: hypothalamus
                                                                                                 32.0%;
                     A; Cross-references: EMBL:X58168
C; Keywords: T-cell receptor
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Best Local Similarity 100.
Matches 2; Conservative
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6 YDKVI 10
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4 QYH 6
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ELAV-like neuronal protein 1, truncated splice form - human
NyAlternate names: Drosophila BLAV(embryonic lethal, abnormal vision)-like 4; Hu antigen
C;Species: Homo sapiens (man)
C;Date: 18-Aug-2000 #sequence_revision 18-Aug-2000 #text_change 09-Jul-2004
C;Accession: 152708
R;Sekido, Y.; Bader, S.A.; Carbone, D.P.; Johnson, B.E.; Minna, J.D.
Cancer Res. 54, 4988-4922, 1994
A;Title: Molecular analysis of the Hub gene encoding a paraneoplastic encephalomyelitis
A;Reference number: 152708; MUID:94349312; PMID:8069866
A;Accession: 152708
A;Accession: 152708
A;Accession: 152708
A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-11 <a href="mailto:sbk">sbk</a>
A;Cross-references: UNIPROT:Q16234; GB:S73887; NID:9688242; PIDN:AAD14142.1; PID:9426184
C;Comment: This abnormal peptide is expressed. For the long splice form, see PIR:138726.
C;Gonment: This abnormal peptide is expressed. For the long splice form, see PIR:138726.
C;Gonetics:
A;Gene: GDB:ELAVL4; HUD; PNEM
A;Cross-references: GDB:141875; OMIM:168360
C;Keywords: alternative splicing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   lipovitellin - African clawed frog
C;Species: Xenopus laevis (African clawed frog)
C;Species: Xenopus laevis (African clawed frog)
C;Date: 18-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 10-Nov-1995
C;Accession: $10624
R;Wallace, R.A.; Hoch, K.L.; Carnevali, O.
J. Mol. Biol. 213, 407-409, 1990
A;Title: Placement of small lipovitellin subunits within the vitellogenin precursor in ?
A;Ritle: Placement $10624; MUID:90278951; PMID:2352275
A;Rocession: $10624
A;Residues: 1-12 <WAL>
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Best Local Similarity 42.9
Matches 3; Conservative
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RILK 9
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S10624
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Search completed: October 21, 2004, 07:27:59 Job time: 16.75 secs

1 KFORPYK 7

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GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
                 Copyright
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OM protein - protein search, using sw model

Run on:

October 21, 2004, 07:05:04; Search time 86 Seconds (without alignments) 60.214 Million cell updates/sec

US-09-720-469A-39 50 Title:

1 KYHRVIKDF 9 Perfect score: Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

1825181 segs, 575374646 residues Searched:

4909 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 12

Post-processing: Minimum Match 08 Maximum Match 1008

Listing first 45 summaries

UniProt 02:* Database :

1: uniprot_sprot:* 2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	O7scd2 neurospora		cydur	anser c	Q8aww0 coscoroba c	Q9xsp2 hylobates s			pygath			saccharc	Q9tqv4 e equus cab		Q9tt31 callithrix		plasn	P83375 serratia pl			Q8mjn2 callithrix	Q8mjn3 callimico g	leontopith	Q8mjn5 saguinus fu		Q8mjn7 saimiri sci	8	Q8mjn9 ateles fusc	Q931x4 vibrio chol	Q8rjfl pseudomonas	25
SUMMARIES	ID	Q7SCD2	Q8AWV7	QBAWV8	Q8AWV9	QBAWWO	Q9XSP2	Q9XSP5	Q9XSP6	Q9XSP7	Q9XSP8	09XSQ4	005403	Q9TQV4	Q9TT29	Q9TT31	Q8N6B1		BS43_SERPL	Q9TWX7	OBMJN1	Q8MJN2	Q8MJN3	Q8MJN4	Q8MJNS	9NJW8Ö	Q8MJN7	OSMJN8	6NLM8Q	Q93LX4	QBRJF1	Q82625
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Q8mel7 sida hooker	Q8mel9 pavonia has	Q8mem2 lagunaria p	Q8mepo hibiscus pe				Q8mer0 hibiscus co		Q8mer7 fioria viti	Q8mer8 dombeya til	Q8mes1 alyogyne pi	Q8mes3 alyogyne cr	Q8mes5 abelmoschus
Q8MEL7	Q8MEL9	Q8MEM2	OBMEPO	Q8MEP3	Q8MEP5	Q8MEQ7	Q8MER0	Q8MER1	Q8MER7	Q8MER8	Q8MES1	QBMES3	Q8MESS
7	7	7	7	~	7	7	~	N	7	~	7	N	7
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## ALIGNMENTS

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A Galagan J.E., Calvo S.E., Borkovich K.A., Selker E.U., Read N.D.,

A Jaffe D., FitzHugh W., Ma L.-J., Smirnov S., Purcell S., Rehman B.,

A Jaffe D., FitzHugh W., Mang S., Nielsen C.B., Butler J., Endrizzi M.,

A Cui D., Ianakiev P., Pedersen D., Nelson M., Washburne M.,

A Selitrennikoff C.P., Kinsey J.A., Braun E.L., Zelter A., Schulte U.,

A Kothe G.O., Jedd G., Mewes W., Staben C., Marcotte E., Greenberg D.,

A Roy A., Poley K., Naylor J., Thomann N., Barrett R., Gnerre S.,

A Ramal M., Kamvysselis M., Mauceli E., Bielke C., Rudd S., Frishman D.,

A Krystofova S., Rasmussen C., Metzenberg R.L., Perkins D.D., Kroken S.,

A Cogoni C., Macino G., Catcheside D., Li W., Pratt R.J., Osmani S.A.,

A DeSouza C.C., Glass L., Orbach M.J., Berglund J., Voelker R.,

A Varden O., Plamann M., Seiler S., Dunlap J., Radford A., Aramayo R.,

A Natvig D.O., Alex L.A., Mannhaupt G., Ebbole D.J., Freitag M.,

RA Natvig D.O., Alex L.A., Mannhaupt G., Ebbole D.J., Freitag M.,

RA Nature O:O-O(2003).

A Nature O:O-O(2003).

- I-CAUTION: The Sequence shown here is derived from an

EMBL/Genbank/DbBJ whole genome shotgun (WGS) entry which is

C BEBL/Genbank/DbBJ whole genome shotgun (WGS) entry which is

EMBL, AABX01000130; EAA34319.1; --

SEQUENCE 9 AA; 1173 MW; C483A32B50440449 CRC64;
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                                                                                                                                                                          Neurospora crassa.
Bukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycetes;
Sordariomycetidae; Sordariales; Sordariaceae; Neurospora.
NCBI_TaxID=5141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
44.0%; Score 22; DB 2; Length 9;
Best Local Similarity 42.9%; Pred. No. 1.8e+06;
Matches 3; Conservative 3; Mismatches 1; Indels
                                                                                                 Last sequence update)
Last annotation update)
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                                       9 AA.
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01-MAR-2004 (TrEMBLrel. 26, Last seq
01-MAR-2004 (TrEMBLrel. 26, Last ann
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                                      PRELIMINARY;
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2 FNRIYKD 8
                                                                                                                                    Predicted protein.
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                                                                                                                                                        Name=NCU05334.1
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01-MAR-2003
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RESULT 1
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8 AA; 911 MW; D73AADC2C05331F1 CRC64;
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Best Local Similarity 50.v.,
'-has 3; Conservative
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Best Local Similarity
Matches 2; Conserv
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2 HKIVAD 7
                                                                                                                                                   3 HRVIKD 8
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Name=LDH-B;
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HKIVAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=8863;
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Q9XSP2
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Anseriformes; Anatidae; Cygnus.
NCBI_TaxID=110926;
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Archosauria, Aves, Neognathae, Anseriformes, Anatidae, Anser.
                                       Anas platyrhynchos (Domestic duck).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi;
Archosauria, Aves, Neognathae, Anseriformes; Anatidae, Anas.
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                                                                                                                 SEQUENCE FROM N.A.
Cotter J.-P., StJohn J., Quinn T.W.;
Submitted (JUN-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AY123327; AAM96901.1; -.
NON_TER 1 1 1
NON_TER 8 8 8 ... n73AADC2C05331F1 CRC64;
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Pred. No. 1.8e+06;
3; Mismatches 1; Indels
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Submitted (JUN-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AY123326; AAM96900.1; -.
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Cotter J.-P., StJohn J., Quinn T.W.;
Submitted (JUN-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AY123325; AAM96899.1; -.
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Last sequence update)
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Best Local Similarity 33.3
Matches 2; Conservative
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2 HKIVAD 7
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         LDH-B (Fragment).
                                                                                                 NCBI TaxID=8839;
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                         Name=LDH-B;
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Q8AWV8
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SEQUENCE FROM N.A.

MEDLINE=20065871; PubMed=10598812;

Bonthron D.T., Smith S.L., Campbell R.;

"Complex patterns of intragenic polymorphism at the PDGFA locus.";

"Complex patterns of intragenic polymorphism at the PDGFA locus.";

Hum. Genet. 105:452-459(1999).

EMBL; AJ243280; CAB45927.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hylobates syndactylus (Siamang) (Symphalangus syndactylus).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hylobatidae, Hylobates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Coscoroba coscoroba (Coscoroba swan).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Archosauria, Aves, Neognathae, Anseriformes, Anatidae, Coscoroba.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        n 42.0%; Score 21; DB 2; Length 8; Similarity 33.3%; Pred. No. 1.8e+06; 2; Conservative 3; Mismatches 1; Indels
                                                          1; Indels
  Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
Cotter J.-P., StJohn J., Quinn T.W.;
Submitted (JUN-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AY123324; AAM96898.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11 AA; 1345 MW; 7FB881F101E1E044 CRC64;
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01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOV-1999 (TrEMBLrel. 19, Last annotation update)
Platelet-derived growth factor A chain (Fragment).
Name=PDGFA;
                                                                                                                                                                                                                                                                                                                                                                          01-MAR-2003 (TrEMBLrel. 23, Created)
01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
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Query Match 42.0%; Score 21; DB 2; I Best Local Similarity 33.3%; Pred. No. 1.8e+06; Matches 2; Conservative 3; Mismatches 1;
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Q9XSP5

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01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Platelet-derived growth factor A chain (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                     11 AA.
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                                                                                Pygathrix nemaeus (Dove langur).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 50.0
Matches 3; Conservative
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6 HHRSVK 11
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6 HHRSVK 11
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                                                                                                                                    NCBI_TaxID=54133;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=98375;
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                                                                                                                         Pygathrix.
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Q9XSQ4;
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EMBL; AJ243277; CAB45926.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
MEDLINE=20065871; PubMed=10598812;
Bonthron D.T., Smith S.L., Campbell R.;
"Complex patterns of intragenic polymorphism at the PDGFA locus.";
Hum. Genet. 105:452-459(1999).
EMBL; AJ243279; CAB45925.1; -.
                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan. 131 TaxID=9598;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pongo pygmaeus (Orangutan).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pongo.
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Pred. No. 3.9e+03;
2; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                       11 AA; 1331 MW; 7FB881F101E1F2D4 CRC64;
                                                                                                       01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Platelet_derived growth factor A chain (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Platelet-derived growth factor A chain (Fragment).
Name=PDGFA;
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                                                                                 11 AA.
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Best Local Similarity 50.0%;
Matches 3; Conservative
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les 3; Conservative
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HHRSVK 11
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                                                                                                                                                              Name=PDGFA;
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SEQUENCE
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                                                                                Q9XSP5
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Q9XSP7
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Q9XSP6
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Mammalia, Butheria, Primates, Catarrhini, Cercopithecidae, Colobinae,
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Mammalia, Eutheria, Primates, Catarrhini, Cercopithecidae, Colobinae,
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MEDLINE=20065871; PubMed=10598812;
Bonthron D.T., Smith S.L., Campbell R.;
"Complex patterns of intragenic polymorphism at the PDGFA locus.";
Hum. Genet. 105:452-459(1999).
EMBL; AJZ43282; CAB45924.1; -.
NON TER 1 1 11
NON TER 11 11
SEQUENCE 11 AA; 1345 MW; 7FB881F101E1E044 CRC64;
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MEDLINE=20065871; PubMed=10598812;
Bonthron D.T., Smith S.L., Campbell R.;
"Complex patterns of intragenic polymorphism at the PDGFA locus.";
Hum. Genet. 105:452-459(1999).
EMBL; AJ243281; CAB46013.1; -.
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01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Platelet-derived growth factor A chain (Fragment).
Name=PDGFA;
01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Platelet-derived growth factor A chain (Fragment).
Name=PDGFA;
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Equus caballus (Horse)
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                                                                                                                                                                                                                                                                                                                                                                                                                 10 AA;
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                                                                    NCBI_TaxID=9796;
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"A 29.425 kb segment on the left arm of yeast chromosome XV contains more than twice as many unknown open reading frames.";
Yeast 11:975-986(1995).
EMBL; X83121; CAA58183.1; -.
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O9TQV4;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
65-JUL-2004 (TrEMBLrel. 27, Last annotation update)
transferrin) (Equus caballus pop-variant Fl transferrin) (Equus caballus pop-variant F2 transferrin) (Equus caballus pop-variant D transferrin) (Equus caballus pop-variant O transferrin) (Equus caballus pop-variant F3 transferrin) (Equus caballus pop-variant F3 transferrin)
                                                                                                                                        SEQUENCE FROM N.A.
MEDLINE=20065871; PubMed=10598812;
Bonthron D.T., Smith S.L., Campbell R.;
"Complex patterns of intragenic polymorphism at the PDGFA locus.";
Hum. Genet. 105:452-459(1999).
EMBL; AJ243278; CAB45916.1; -.
                         Gorilla gorilla (gorilla).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Gorilla.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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Q05403;
Q1-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
Orf 00915 protein (Fragment).
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MEDLINE=96021609; PubMed=8533473;
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HNVVK 7
                                                                                                 NCBI_TaxID=9593;
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         Name=PDGFA;
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Q05403
ID Q0540
AC Q0540
DT 01-NO
DT 0
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Matches
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Q9TQV4
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O9TT31;
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 13, Last annotation update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Beta-globin (Fragment).
Callithrix jacchus (Common marmoset).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Platyrrhini; Callitrichidae; Callithrix.
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Platyrrhini; Callitrichidae; Saguinus.
Eŭkaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Perissodactyla, Equidae, Equus.
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MEDLINE=20188801; PubMed=10723742;
Francino M.P., Ochman H.;
"Strand symmetry around the beta-globin origin of replication in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 20; DB 2; Length 10;
Pred. No. 5.5e+03;
1; Mismatches 0; Indels
                                                                                                                               Bell T.K.;
EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                 6DF67DE9D1AEBDDB CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Beta-globin (Fragment).
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Mol. Biol. Evol. 17:416-422(2000).
EMBL; AF205415; AAF23766.1; -.
                                                                                                                                                                                                       EMBL, AF185770; AAF05483.1;
EMBL, AF185771; AAF05484.1; --
EMBL; AF185772; AAF05485.1; --
EMBL; AF185773; AAF05486.1; --
EMBL; AF185774; AAF05487.1; --
EMBL; AF185775; AAF05488.1; --
EMBL; AF185775; AAF05489.1; --
EMBL; AF185776; AAF05489.1; --
                                                                                                        SEQUENCE FROM N.A.
Giffard J.M., Brandon R.B.,
Submitted (SEP-1999) to the
EMBL; AF185778; AAF05491.1;
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Best Local Similarity 75.v°,
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          TISSUE=Blood;
MEDLINE=20188801; PubMed=10723742;
Francino M.P., Ochman H.;
"Strand symmetry around the beta-globin origin of replication in primates.";
Mol. Biol. Evol. 17:416-422(2000).
EMBL; AF205413; AAF23764.1; -.
NON TER 1 1 1
SEQUENCE 12 AA; 1251 MW; 07BC8AFAE8D72DD4 CRC64;
                                                                                                                                                                              Query Match 40.0%; Score 20; DB 2; Length 12; Best Local Similarity 100.0%; Pred. No. 6.6e+03; Matches 3; Conservative 0; Mismatches 0; Indels
NCBI_TaxID=9483;
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                                                                                                        October 21, 2004, 07:04:19 ; Search time 83 Seconds (without alignments) 38.898 Million cell updates/sec
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Adc17702 C
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Aar91898 F
Aar45361 C
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GenCore version 5.1.6
(c) 1993 - 2004 Compugen Ltd.
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ABR16875
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ADC17702
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Gapop 10.0 , Gapext 0.5
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geneseqp2001s:*
geneseqp2002s:*
geneseqp2003as:*
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length: 12
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Abr16761 Human can	Abr16123 Human can	Abr16487 Human can	Abr17152 Human can	Abr16466 Human can	Abr16141 Human can	Abr16299 Human can	Abr16898 Human can	Aar72938 E. coli P	Aar72894 E. coli P		Adp49297 Natriuret	Aar72949 E. coli P	Aar72905 E. coli P	Aam42948 Mycoplasm	Aaw21441 HSV glyco	Aao31067 Human F8	Aar20729 Pancreas-	Aay51905 Elastase	Aam22987 HIV pepti
ABR16761	ABR16123	ABR16487	ABR17152	ABR16466	ABR16141	ABR16299	ABR16898	AAR72938	AAR72894	ADP49294	ADP49297	AAR72949	AAR72905	AAM42948	AAW21441	AA031067	AAR20729	AAY51905	AAM22987
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27			27	27	27	27	27	56	56	56	26	26	56	26	26	56	25	25	25
26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	4.57

### ALIGNMENTS

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This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                              Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                               Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
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                                                                                        Human cyclophilin B peptide fragment #43
                                                                                                                                                                                                                                                                                                                                                                                                      Claim 10; Page 60; 64pp; Japanese.
              AAY69963 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                             (SUMU ) SUMITOMO PHARM CO LID. (ITOH/) ITOH K.
                                                                                                                                                                                                                            99WO-JP003360.
                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                          WPI; 2000-116932/10.
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Best Local Similarity
Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 9 AA;
                                                                                                                                                                            WO9967288-A1.
                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                            24-JUN-1999;
                                                                                                                                                                                                                                                     25-JUN-1998;
                                                              11-APR-2000
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                                      AAY69963;
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AAY69963
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                capable of expressing the interferons and/or antigenic proteins (AP), antigenic peptides derived from the proteins or DNAs capable of expressing the antigenic proteins or peptides, in the manufacture of an agent for induction of antigen-specific T cells. The products of the invention have virucide and cytostatic activity and can be used for gene therapy or as inducers of antigen-specific T cells. The action of interferon-alpha (IFN-alpha) in a system for inducing specific cytotoxic T cell (CTL) by administering an antigenic peptide in an incomplete Freund's adjuvant (IFA) emulsion preparation form was evaluated. IFNS (or inducing antigen-specific T cells in an individual who has been administered with AP (or DNA encoding AP) or vice versa. The medicament is useful for the treatment or prophylaxis of a tumor or a viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Use of antigenic proteins, peptides, interferon or their encoding DNA, in the manufacture of an agent for the induction of antigen-specific T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes the novel use of interferons (IFNs) or DNAs
                                                                                                                                                            Tumor antigenic peptide; interferon; IFN; antigen-specific T cell; virucide; cytostatic; gene therapy; IFN-alpha; cytotoxic T cell; viral infectious disease; cyclophilin B; human.
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                                                                                                                               Human cyclophilin B derived tumor antigenic peptide SEQ ID 33.
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                                  AAB46947 standard; peptide; 9 AA.
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                                                                                                                                                                                                                                                                                                                                                                                         (SUMU ) SUMITOMO PHARM CO LID
                                                                                                                                                                                                                                                                                                                         24-JUL-2000; 2000EP-00306263.
                                                                                                                                                                                                                                                                                                                                                          99JP-00207687.
                                                                                               (first entry)
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Matches 9; Conser
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                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                          22-JUL-1999;
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                                                                 AAB46947;
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                                              suppressor; antigen;
desensitization;
                                                                                                                                                                                                                                                                                                                                                                          Desensitizers or allergic reaction suppressors containing peptides originating from the same antigenic substances as reaction-inducing substances, useful for preventing or treating type I allergic diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 51; DB 7; Length 9; 100.0%; Pred. No. 1.7e+06; tive 0; Mismatches 0; Indels
                                     cytostatic; antiallergic; vaccine; allergic reaction allergy; epitope; cancer; immunoglobulin B antibody; cyclophilin B.
             Modified cyclophilin B protein amino acids 91-99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 8; SEQ ID NO 9; 49pp; Japanese.
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                                                                                                                                                                                                            28-AUG-2002; 2002WO-JP008641.
                                                                                                                                                                                                                                            29-AUG-2001; 2001JP-00260046
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                                                                                                                                                                                                                                                                                                                                                                                                                             and in cancer vaccines
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                                                                                                                                                                                                                                                                                                          Itoh K, Yamada A;
                                                                                                                                             WO2003020306-A1.
                                                                                                                                                                                                                                                                              (ITOH/) ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 9 AA;
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                                                                                                                                                                             13-MAR-2003
                                                                                                              Synthetic.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes the novel use of interferons (IFNs) or DNAs capable of expressing the interferons and/or antigenic proteins (AP), antigenic peptides derived from the proteins or DNAs capable of expressing the antigenic proteins or peptides, in the manufacture of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Use of antigenic proteins, peptides, interferon or their encoding DNA, the manufacture of an agent for the induction of antigen-specific T
                                                                                                                                                 This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                Gaps
                                                                                                    Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                 Tumor antigenic peptide, interferon, IFN; antigen-specific T cell; virucide, cytostatic; gene therapy; IFN-alpha; cytotoxic T cell; viral infectious disease; cyclophilin B; human.
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                                                                                                                                                                                                                                                                                                                                                                 Human cyclophilin B derived tumor antigenic peptide SEQ ID 17.
                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                              Length 9;
                                                                                                                                                                                                            Score 47; DB 3; 1
Pred. No. 1.7e+06;
1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 15; 25pp; English.
                                                                                                                                                                                                                                                                                                            AAB46931 standard; peptide; 9 AA.
                                                                                                                                Claim 4; Page 49; 64pp; Japanese
                                                                                                                                                                                                                             1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yamaoka T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SUMU ) SUMITOMO PHARM CO LID
                                         CO LTD
       99WO-JP003360,
                       98JP-00178449.
                                                                                                                                                                                                             92.2%;
88.9%;
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                                         (SUMU ) SUMITOMO PHARM
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Best Local Similarity
8; Conserve
                                                                                     WPI; 2000-116932/10.
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                                                                                                                                                                                                                                                 1 DYMIQGGDF
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                                                                   Gomi S;
                                                                                                                                                                                             Sequence 9 AA;
                                                   ITOH
       24-JUN-1999;
                        25-JUN-1998;
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                                                                                                                                                                                                                                                                                                                              AAB46931;
                                                   (ITOH/)
                                                                   toh K,
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The present invention describes peptides recognising cancer cells with tumour specific cytotoxic T lymphocytes (CTD). The peptides recognising cancer cells with tumour specific CTLs are selected from: (1) peptides of sequences (AAG68066 to AAG68069); (2) peptides containing the above mentioned sequences; (3) peptides having 70 % or more of homogeneity with the above mentioned sequences; and (4) peptides with one or more deleted, substituted, added or inserted amino acid(s) of the above mentioned sequences, particularly those having recognising property due to HLA-A2402 binding CTL, especially having at least 5 amino acids, used for medicine, particularly anticancer agents, derived from antitumour antigenic peptides of lck, src family, SART-1, SART-3 or cyclophilin B genes. The antitumour peptides have cytostatic activities. The peptides are used for the treatment of cancer. The peptides cause activation of the contained from the c
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agent for induction of antigen-specific T cells. The products of the invention have virucide and cytostatic activity and can be used for gene therapy or as inducers of antigen-specific T cells. The action of interferon-alpha (IFN-alpha) in a system for inducing specific cytotoxic T cell (CTL) by administering an antigenic peptide in an incomplete Freund's adjuvant (IFA) emulsion preparation form was evaluated. IFNs (or DNA encoding IFNs) are useful in the manufacture of a medicament for inducing antigen-specific T cells in an individual who has been administered with AP (or DNA encoding AP) or vice versa. The medicament is useful for the treatment or prophylaxis of a tumor or a viral
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antitumour; cancer; cancer cell redognition; antigenic; CTL; lck; src; tumour specific cytotoxic I lymphocyte; anticancer; SART-1; SART-3; cyclophilin B gene; HLA-A2402.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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Pred. No. 1.7e+06;
1; Mismatches 0;
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88.9%;
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Best Local Similarity 88.9
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           infectious disease
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1 DFMIQGGDF
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Sequence 9 AA;

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The invention relates to a composition (I) comprising an immune effector cell and a cell penetrating peptide (CPP) associated with an antigen or antibody. Also included are (1) a vaccine comprising (I), CPP associated with an antigen, and apharmaceutically acceptable carrier and (2) preparing a composition for a disease, by providing (I) and CPP associated with an antigen for disease, and introducing the antigen associated CPP to (I), where antigen enters into the cell. The antigens care, for example, tumour antigen derived epitopes recognised by tumour infiltrating lymphocytes (TIL) of Him human leukocyte antigen) class I or II. The composition is useful for enhancing immunity in an animal to a disease, by administering a mature dendritic cell comprising CPP congrises both CD4+ and CD8+ T cells. It is also useful for treating a sociated with an antigen to disease, to the animal, such that following cancer, non-Hodgkin's lymphoma, lymphoma, uterine cancer, non-Hodgkin's lymphoma, leukaemia, lymphoma, uterine cancer, non-Hodgkin's lymphoma, uterine cancer, prostate cancer, kidney cancer, adenocarcinoma, breast cancer, prostate cancer, vorarian cancer and pancreatic cancer). The animal is further subjected to a cancer treatment including surgery, breast cancer, prostate cancer, vorarian cancer and pancreatic cancer). The animal is further subjected to a cancer treatment including surgery, cradiation, chemotherapy or gene therapy. The administration of (I), preferably dendritic cell is prior to, subsequent to or concurrent with, the cancer treatment. The present sequence is a tumour antigen derived ceptived epitope for inclusion in the composition of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                       Cell penetrating peptide; cancer; tumour; melanoma; thymoma; antigen;
                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                   lymphoma; sarcoma; lung cancer; non-Hodgkin's lymphoma; leukaemia; Hodgkin's lymphoma; leukaemia; Hodgkin's lymphoma; leukaemia; kidney cancer; adenocarcinoma; breast cancer; prostate cancer; ovarian cancer; pancreatic cancer; epitope; vaccine; dendritic cell; tumour infiltrating lymphocyte; TIL; human leukocyte antigen; HLA; cytostatic; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel composition for treating a disease in an animal, comprises an immune effector cell and cell penetrating peptide associated with an antigen or antibody.
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                                                                                                                                                                                                                                                                                                                                                                             Human Cyp-B class I HLA widely expressed antigen peptide #2.
Score 47; DB 4; Length 9;
Pred. No. 1.7e+06;
1; Mismatches 0; Indels
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                                                                                                                                                                                                                                              ABG79076 standard; peptide; 9 AA.
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   92.2%;
                                                                                                                                                                                                                                                                                                                                    (first entry)
                                               8; Conservative
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DFMIQGGDF
                                                                                           1 DYMIQGGDF
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                 Antigen specific T-cell; detection; diagnosis; cancer specific T-cell; cancer; tumour; cervical cancer; prostate cancer; cellular immunity; immune therapy; cytostatic; immunostimulant; vaccine; antigenic peptide; human; human leukocyte antigen; HLA-A24 epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A detection method of antigen specific T-cells, comprises the use of plural antigenic peptides, useful in semi-quantitative determination of cancer specific T-cell frequencies and for monitoring cellular immunity.
                                                                                   Gaps
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                                             Score 47; DB 5; Length 9; Pred. No. 1.7e+06; 1; Mismatches 0; Indels
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                                             92.2%;
nilarity 88.9%;
Conservative 1
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              Sequence 9 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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Best Local Similarity
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                                  WO2003100027-A2
                                                                                                                                                                                                                                                                                                                         Sequence 9 AA;
              Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-JUN-1998;
                                                       04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                   ..
60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY69958;
                                                                                                                                                                                                                                                                                                      antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Itoh K,
                                                                                                                                          Wang R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ITOH/)
                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY69958
    a
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                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to allergic reaction suppressors, comprising a peptide which originates from the same antigenic substance as the antigenic substance inducing the (peptide-originated) allergic reaction, and containing an epitope different from the epitope participating in the induction of the allergic reaction and yet dos not induce the allergic reaction. The desensitizers or allergic reaction suppressors are useful for treating and preventing type I allergic diseases and in cancer vaccines for preventing or treating cancer. The allergic reaction or suppressors are also useful for suppressing an allergic reaction or reducing immunoglobulin E antibody production, and for desensitization to antigens. This sequence represents a peptide used in the invention and corresponds to amino acids 91-99 of the cyclophilin B protein.
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0
                                                                                                      vaccine; allergic reaction suppressor; antigen;
immunoglobulin E antibody; desensitization;
                                                                                                                                                                                                                                                                                                                    Desensitizers or allergic reaction suppressors containing peptides originating from the same antigenic substances as reaction-inducing substances, useful for preventing or treating type I allergic diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              metastatic cancer cell differentiation; mutated fibronectin; metastatic cancer; class I HLA-restricted; widely antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 47; DB 7; Length 9;
Pred. No. 1.7e+06;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Class I HLA-restricted widely expressed antigen #7.
                                                                                  Cyclophilin B protein amino acids 91-99.
                                                                                                                                                                                                                                                                                                                                                                        Claim 8; SEQ ID NO 2; 49pp; Japanese.
                    ADC17702 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADG89642 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1;
                                                                                                                                                                                                             28-AUG-2002; 2002WO-JP008641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92.2%;
                                                                                                                                                                                                                                  29-AUG-2001; 2001JP-00260046
                                                             (first entry
                                                                                                      ic; antiallergic;
epitope; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8; Conservative
                                                                                                                                                                                                                                                                                                                                                     and in cancer vaccines
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 DYMIQGGDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 DFMIQGGDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                           Itoh K, Yamada A;
                                                                                                                                                                     WO2003020306-A1
                                                                                                                 allergy; epitc
cyclophilin B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9 AA;
                                                                                                                                                                                                                                                      HOTI (/HOTI)
                                                             18-DEC-2003
                                                                                                      cytostatic;
                                                                                                                                                                                          13-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-MAR-2004
                                                                                                                                                Synthetic.
                                         ADC17702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG89642;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 10
RESULT 9
            ADC17702
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Identifying a cell that differentiates into a metastatic cancer cell, useful for preventing metastatic cancer, comprises identifying a mutated fibronectin in the cell.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention comprises a method for identifying a cell that will differentiate into a metastatic cancer cell, the method involves identifying a mutated fibronectin in the cell. The method of the invention is useful for preventing metastatic cancer. The present a acid sequence represents a Class I HLA-restricted widely expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 47; DB 8; Length y;
Pred, No. 1.7e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO 85; 137pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human cyclophilin B peptide fragment #38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY69958 standard; peptide; 9 AA
                                                                                                                             (BAYU ) BAYLOR COLLEGE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SUMU ) SUMITOMO PHARM CO LTD
28-MAY-2003; 2003WO-US016736.
                                                                 28-MAY-2002; 2002US-0383530P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92.2%;
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Matches
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                                 RESULT 1.
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                                  This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL;
HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
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                                                                                                                                                                                                Score 41; DB 3; Length 9; Pred. No. 1.7e+06;
                                                                                                                                                                                                                                          o; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Trp, Met
                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human cyclophilin B peptide fragment #41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /label= Phe, Leu, Ile,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           label= Phe, Tyr, Met,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 10; Page 60; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY69961 standard; peptide; 9 AA.
Claim 4; Page 61; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SUMU ) SUMITOMO PHARM CO LTD.
                                                                                                                                                                                                    80.4%;
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Best Local Similarity 87.5
Fra 7; Conservative
                                                                                                                                                                                                                          Best Local Similarity 87.5
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-116932/10
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|DFMIQGGD 8
                                                                                                                                                                                                                                                                                            DYMIQGGD 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9 AA;
                                                                                                                                                          Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ITOH/) ITOH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9967288-A1
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                                                                                                                                                                                                      Query Match
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Keratinocyte growth factor; KGF; Cysl; Cysl; deletion; substitution; Met-KGF analog; stimulate; non-fibroblast epithelial cell; antibody; detection; purification; adnexal structure; hair follicle; sweat gland; sebaceous gland; burn; gastric ulcer; duodenal ulcer; liver function; hepatic cirrhosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                              Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             70.6%; Score 36; DB 3; I
75.0%; Pred. No. 1.7e+06;
tive 1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KGF-b minor component N-terminal fragment.
                                                                                                                 Human cyclophilin B peptide fragment #37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR91898 standard; protein; 12 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; Page 61; 64pp; Japanese.
AAY69957 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                           (SUMU ) SUMITOMO PHARM CO LTD
                                                                                                                                                                                                                                                                                                                                99WO-JP003360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-DEC-1996 (first entry)
                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-116932/10.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gomi S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 9 AA;
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                                                                                                                                                                                                                                                                                                                                24-JUN-1999;
                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                      WO9967288-A1
                                                                                                                                                                                                                                                                                                                                                                      25-JUN-1998;
                                                                             11-APR-2000
                                                                                                                                                                                                                                                                                          29-DEC-1999
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                                     AAY69957;
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1 DYMIQGGD 8

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The sequences given in AAR91897-98 represent the N-terminal peptides from the major and minor components, KGF-a and KGF-b, of recombinantly capressed keratinocyte growth factor (KGF). Interestingly the N-terminal fragment of KGF-b indicates that it is an N-terminally truncated form of KGF which is proteolytically cleaved at the Arg23-Ser24 peptide bond. KGF may be used in the construction of KGF analogues (see also AAR91865-96).

These analogues have up to the first 24 N-terminal amino acids modified so that Cys1 and Cys15 are deleted or substituted. Deleting or substituting the Cys residues at positions 1 and 15 of native KGF substituting the Cys residues at bositions 1 and 15 of native KGF produces a KGF analogue which has substantially improved stability. These Expensions of native KGF analogues for the detection of non-fibroblast epithelium agents and in the generation of antibodies for the detection or purification of native KGF. The KGF analogues may be applied to tissues with damage to, or insufficient numbers of, non-fibroblast epithelium cells. This includes the stimulation, proliferation and differentiation of adnexal structures, e.g. hair follicles, sweat glands, and sebaceous glands in burns patients, treating gastric and duodenal ulcers, and increasing liver function to treat or prevent e.g. hepatic cirrhosis
                                                                                                                                                                                                                                                           Keratinocyte growth factor analogues with modified N-terminal region stimulate prodn. of non-fibroblast epithelial cells and are useful e.g. to treat gastric and duodenal ulcers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Preparation; high; purity; yield; chemical analysis; esterase; determination; phospholipid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 27.5; DB 2; Length 12; Pred. No. 2.4e+02; 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                    Example 3; Page 52; 118pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR45361 standard; peptide; 9 AA.
                                                                                                                                                                                            Chen B,
                                                                                 94US-00323340.
94US-00323475.
95US-00487825.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                53.9%;
75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93JP-00008835
                                                  95WO-IB000971.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Saccharomonospora glauca
                                                                                                                                                                                            Morris CF, Kenney WC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                        (AMGE-) AMGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 12 AA;
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04-JUL-1994
                                                  12-OCT-1995;
                                                                                    13-OCT-1994;
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                                                                                                    13-OCT-1994;
                                                                                                                     07-JUN-1995;
                 25-APR-1996
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Job time : 84 secs

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DNA fragment for choline oxidase prepn., for high purity and high yield comprises gene coding choline oxidase prepd. by culturing microbe transformed by recombinant plasmid for choline chemical analysis and
                                                                                                                                                                                                          The sequence is that of the N-terminal of choline oxidase. Choline oxidase is useful for chemical analysis of choline, determination of choline esterase contained in serum and determination of phospholipid. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                       ..
                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                     Score 27; DB 2; Length 9; Pred. No. 1.7e+06; 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Search completed: October 21, 2004, 07:21:04
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                                                                                                                                                                                 Claim 5; Page 2; 8pp; Japanese.
            92JP-00009256
                                                                                                                                                                                                                                                                                                                       52.9%;
                                          (TAKE ) TAKEDA CHEM IND LTD.
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                                                                       WPI; 1994-011021/02
                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
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DYVVVGG 9
                                                                                                                                                                                                                                                                                                                                                                                   1 DYMIQGG 7
                                                                                                                                                                                                                                                                                            Sequence 9 AA;
            22-JAN-1992;
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; CELL TYPE: pancreatic US-08-457-172-1
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                                                                                                                                                                                                                             US-08-457-172-1
Sequence 1, Appli
Sequence 75, Appl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 8, Appli
Patent No. 5196404
Patent No. 5433940
Sequence 1, Appli
Sequence 19, Appli
                                                                                                                                                                                                                                                                                                                                                    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                               October 21, 2004, 07:13:10; Search time 21.5 Seconds (without alignments) 27.761 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                     Description
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1
Sequence 6
Sequence 2
Sequence 2
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                                                                                                                                                                                                                                                                          Issued_Patents_AA:*
1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
         GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-457-172-1

US-08-406-330-75

US-08-556-597-75

US-08-459-568-75

US-08-399-411-75

US-08-516-859A-75

US-09-586-472-75

US-09-586-472-75

US-09-586-472-75

US-09-586-472-75

US-09-586-472-75

US-09-586-472-75

US-09-846-350A-1

US-07-620-426B-19

US-07-620-426B-19

US-07-620-426B-19

US-07-620-426B-19

US-07-620-426B-19

US-07-620-426B-19

US-07-620-426B-19

US-07-620-426B-19

US-08-68-130-21

US-08-68-130-21

US-09-315-444-51

US-09-315-444-51

US-09-721-362-51
                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-841-636A-27
                                                                                                                                                                      478139 segs, 66318000 residues
                                                                                                                                                                                                                                     Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                              using sw model
                                                                                                                                           BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                  US-09-720-469A-40
51
                                                                                                                        1 DYMIQGGDF 9
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Match Length
                                              - protein search,
                                                                                                                                                                                                           Minimum DB seq length: 0 Maximum DB seq length: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                        Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                     Score
                                                                                                                                           Scoring table:
                                                OM protein
                                                                                                                         Sequence:
                                                                                                                                                                      Searched:
                                                                                                                                                                                                                                                                            Database
                                                                  Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Result
                                                                                                                                                                                                                                                                                                                                                                                                                     No.
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Sequence 1, Application US/08457172
; Sequence 1, Application US/08457172
; Patent No. 5622837
; GENERATION:
; GENERATION:
; APPLICANT: Hans Scheefers, Ursula Scheefers-Borchel and
APPLICANT: Hans Scheefers, Ursula Scheefers-Borchel and
TITLE OF INVENTION: DANCREAS ELASTASE 1-SPECIFIC ANTIBODY, A PROCESS FOR
TITLE OF INVENTION: OBTAINING IT, AND A TEST KIT CONTAINING SUCH ANTIBODY
; NUMBER OF SEQUENCES:
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Cooper & Dunham
; STREET: 30 Rockefeller Plaza
; CITY: New York
; STATE: New York
; COUNTRY: USA
; ZIP: 10112
                          58, Appl
444, App
444, App
23, Appl
15, Appl
13, Appl
17, Appl
                                                                                                                                                                                                  Sequence
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Sequence
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Sequence
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                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                               Sequence
US-08-215-805A-52
US-08-215-805A-58
US-08-637-759B-444
US-08-871-355A-444
US-09-201-945-444
US-09-964-201A-23
US-09-964-201A-13
US-08-082-849B-15
US-09-785-921A-13
PCT-US94-01624-15
US-08-410-941-17
US-08-468-546-17
US-08-468-547-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRATIONS SISTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/457,172
FILING DATE:
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/220,395
FILING DATE:
APPLICATION NUMBER: US 07/969,173
FILING DATE:
APPLICATION NUMBER: US 07/969,173
FILING DATE:
APPLICATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 42883/JPW/AJM
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEFAX: (212) 977-9550
TELEFX: 422523 COOP UI
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                  ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   unknown
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MOLECULE TYPE: protein
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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us-09-720-469a-40.rai

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   Score 25; DB 1; Length 9; Pred. No. 3.8e+05;
                                                                                                                                                                                                        Sequence 75, Application US/08406330
Patent No. 5817748
GENERAL INFORMATION:
APPLICANT: Miller, Jonathan L.
TITLE OF INVENTION: MIMOTOPES AND ANTI-MIMOTOPES OF
TITLE OF INVENTION: HUMAN PLATELET GLYCOPROTEIN ID/IX
NUMBER OF SEQUENCES: 81
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon, Hargrave, Devans & Doyle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 75, Application US/08556597
Patent No. 5877155
GENERAL INFORMATION:
APPLICANT: Miller, Jonathan L.
APPLICANT: Lyle, Vicki A.
TITLE OF INVENTION: MIMOTOPES AND ANTI-MIMOTOPES OF
TITLE OF INVENTION: HUMAN PLATELET GLYCOPROTEIN ID/IX
NUMBER OF SEQUENCES: 173
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon, Hargrave, Devans & Doyle LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 14603

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/406,330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      49.0%; Score 25; DB 2; I 37.5%; Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                                                 STREET: Clinton Square, Devans & Doyle STREET: Clinton Square, P.O. Box 1051 CITY: Rochester STATE: New York COUNTRY: USA
                                       1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Timian, Susan J.
REGISTRATION NUMBER: 34,103
REFERENCE/DOCKET NUMBER: 20884/100
TELECOMMUNICATION INFORMATION:
TELEPHONE: (716) 263-1636
TELEFAX: (716) 263-1600
INFORMATION FOR SEQ ID NO: 75:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
49.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 37.5
                   Best Local Similarity 66.7
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; MOLECULE TYPE: peptide US-08-406-330-75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ::::|| |
FVVRGGTF 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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| MVAGGD 7
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CLASSIFICATION:
                                                                            3 MIQGGD 8
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US-08-406-330-75
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US-08-556-597-75
   Query Match
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Patent No. 5811304

GENERAL INFORMATION:

APPLICANT: Huang, Shi

TITLE OF INVENTION: Zinc Finger Proteins

NUMBER OF SEQUENCES: 93

CORRESPONDENCE ADDRESS:

ADDRESSEE: Campbell and Flores

STREET: 4370 La Jolla Village Drive, Suite 700

CITY: San Diego

STATE: California

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        49.0%; Score 25; DB 2; Length 12; 37.5%; Pred. No. 1.6e+02; Live 4; Mismatches 1; Indels
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COMPUTER READBLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/459,568
FILING DATE: 02-JUN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/399,411
FILING DATE: 06-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.3C
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/556,597
Clinton Square, P.O. Box 1051
                                                                                                                                                                                                                                                                                                                                       FILING DATE: 17-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Timian, Susan J.
REGISTRATION NUMBER: 34,103
REFERENCE/DOCKET NUMBER: 20884/101
TELECOMMUNICATION INFORMATION:
TELEPHONE: (716) 263-1636
TELEFAX: (716) 263-1600
INFORMATION FOR SEQ ID NO: 75:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                       FILING DATE:
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/406,330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Campbell, Cathryn A. REGISTRATION NUMBER: 31,815
                                                                               ZIP: 14603
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 3/...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 YMIQGGDF 9
                        Rochester
                                           New York
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1 FVVRGGTF
                                                                USA
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                                                                COUNTRY:
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RESULT 7
US-09-586-472-75
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APPLICANT: Huang, Shi
TITLE OF INVENTION: Retinoblastoma Protein - Interacting
TITLE OF INVENTION: Zinc Finger Proteins
NUMBER OF SEQUENCES: 93
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 24; DB 2; Lengtn 0;
Pred. No. 3.8e+05;
                                                                                                                                                                                    Score 24; DB 2; Length 8; Pred. No. 3.8e+05; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COPERATING SYSTEM: FC-DOS/FGS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/399,411
FILING DATE: 06-MAR-1995
CLASSIFICATION: 06-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 1264
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-8949
INFORMATION FOR SEQ ID NO: 75:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
CITY: San Diego
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                           3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 6
US-08-516-859A-75
; Sequence 75, Application US/08516859A
; Patent No. 6069231
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               Sequence 75, Application US/08399411 Patent No. 5831008
             TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 75:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                      47.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            47.1%;
                                                                                                                                                                    Query Match
Best Local Similarity 50.00,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 50.07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 8 amino acids
                                                                                                                   TYPE: amino acid;
TOPOLOGY: linear
US-08-459-568-75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
                                                                                                                                                                                                                                                                              :::|||
3 LLEGGD 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    :::|||
3 LLEGGD 8
                                                                                                                                                                                                                                                           3 MIQGGD 8
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US-08-399-411-75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 24; DB 3; Length 8; Pred. No. 3.8e+05; 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
CITY: San Diego
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/586,472
FILING DATE: 01-Jun-2000
                                                                                                                                                                                                             CUMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/516,859A
FILING DATE: 18-AUG-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/399,411
FILING DATE: 06-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/292,683
FILING DATE: 18-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAMB: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: 9-LJ 1776
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-9001
TELEFAX: (619) 535-8049
INFORMATION FOR SEQ ID NO: 75:
CENTEMATION CATHRORY
                                                                                                              ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3; Mismatches
APPLICANT: Huang, Shi
TITLE OF INVENTION: Retinoblastoma Protes
TITLE OF INVENTION: Zinc Finger Proteins
NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 75, Application US/09586472
Patent No. 6323335
GENERAL INFORMATION:
APPLICANT: Huang, Shi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 92122
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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Best Local Similarity 50.0.
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                                                                                                                                                      CITY: San Diego
STATE: California
COUNTRY: USA
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SEQUENCE CHARACTERISTICS:
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Best Local Similarity
Matches 4; Conservat
                                          ; TYPE: amino acid
; TOPOLOGY: linear
US-09-528-706-75
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STREET; SUL
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USA
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Sequence 75, Application US/09528706

Sequence 75, Application US/09528706

Patent No. 6468985

GENERAL INFORMATION:

APPLICANT: Huang, Shi

TITLE OF INVENTION: Zinc Finger Protein - Interacting

TITLE OF INVENTION: Zinc Finger Proteins

NUMBER OF SEQUENCES: 106

CORRESPONDENCE ADDRESS:

ADDRESSEB: Campbell & Flores LLP

STREET: 4370 La Jolla Village Drive, Suite 700

CITY: San Diego

STATE: California

COUNTRY: USA

ZIP: 92122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              47.1%; Score 24; DB 3; Length 8; 50.0%; Pred. No. 3.8e+05; tive 3; Mismatches 0; Indels
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               PRIOR APPLICATION DATA:

APPLICATION DATA:

APPLICATION NUMBER: US 09/528,706

FILING DATE: 17-MAR-2000

APPLICATION NUMBER: US 08/516,859

FILING DATE: 18-AUG-1995

APPLICATION NUMBER: US 08/399,411

FILING DATE: 06-MAR-1995

APPLICATION NUMBER: US 08/292,683

FILING DATE: 18-AUG-1994

ATTORNEY/AGENT INFORMATION:

REFERENCE/DOCKET NUMBER: P-LJ 4130

TELECOMMUNICATION INFORMATION:

TELEPHONE: (619) 535-9001

TELEPHONE: (619) 535-9001

TELEFAX: (619) 535-9001

TELEFAX: (619) 535-9001

TELEFAX: (619) 535-9001
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PRIOR APPLICATION NUMBER: US 08/292,683

FILING DATE: 18-AUG-1994

ATTORNEY/AGBNT INFORMATION:

NAME: Campbell, Cathryn A.

REGISTRATION NUMBER: 31,815

REFERENCE/DOCKET NUMBER: P-LJ 1776

TELECHONE: (619) 535-9001

TELEPHONE: (619) 535-8949

INFORMATION FOR SEQ ID NO: 75:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE DESCRIPTION: SEQ ID NO: 75:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/516,859
FILING DATE:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
     CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                     LENGIH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
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Best Local Similarity
Matches 3; Conserval
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3 LLEGGD 8
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STREET: 500 West Madison Street, Suite 3800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       h 47.1%; Score 24; DB 3; Length 10; Similarity 100.0%; Pred. No. 2e+02; 4; Conservative 0; Mismatches 0; Indels
                                             0; Indels
Length 8;
                                                                                                                                                                                                                                                                                                                         APPLICANT: DINKELBORG, LUDGER
APPLICANT: STEPHAN, CHRISTOPH
APPLICANT: SEMMLER, WOLFHARD
APPLICANT: SPECK, ULRICH
APPLICANT: KENKLEIN, PETER
TITLE OF INVENTION: COMPLEX COMPOUNDS FOR DIAGNOSIS OF
TITLE OF INVENTION: VASCULAR DISEASES
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
47.1%; Score 24; DB 4; 1
50.0%; Pred. No. 3.8e+05;
tive 3; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: USA
ZIP: 60661-2511
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: COMPAQ Prolinea 4/25s
OPERATING SYSTEM: DOS
SOFTWARE: WordPerfect for Windows 6.0a
APPLICATION NUMBER: US/08/765,953
FILING DATE: 17-JULY-1997
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DE95/00837
FILING DATE: 13-JULY-1995
APPLICATION NUMBER: P 44 25 778.3
FILING DATE: 13-JULY-1994
ATTORNEY/AGENT INFORMATION:
NAME: WORTIMER: JOHN S.
REGISTRATION NUMBER: 30,407
REFERENCE/DOCKET NUMBER: 1214.00001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/876-1800
TELEFAX: 312/876-2020
                                                                                                                                                                                                                                                          Sequence 8, Application US/08765953; Patent No. 6342201; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
 Query Match
Best Local Similarity 50.0
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: peptide
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                                                                                                                                                                        Length 12;
                                                                                                                                                                                                                                 i; Indels
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 720 KB storage
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Macintosh 6.0.7
SOFTWARE: MicrosoftWord 4.0
                                                                                                                                                                   Query Match 47.1%; Score 24; DB 4; I Best Local Similarity 42.9%; Pred. No. 2.5e+02; Matches 3; Conservative 3; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19, Application US/07620426B
Sequence 19, Application US/07620426B
Patent No. 5298421
GENERAL INFORMATION:
APPLICANT: Davies, Huw Maelor
APPLICANT: Pollard, Michael Roman
APPLICANT: Voekler, Toni Alois
TITLE OF INVENTION: Plant Thioesterases
NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/07/620,426B
FILING DATE: 19901130
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/514,030
FILING DATE: 26-APRIL-1990
ATTORNEY/AGENT INFORMATION:
NAME: Blizabeth Lassen
REGISTRATION NUMBER: 31,845
NAME: Donna E. Scherer
REGISTRATION NUMBER: 34,719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CGNE 70-1
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US-07-662-007B-19
; Sequence 19, Application US/07662007B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Plant Thic
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Calgene, Inc.
STREET: 1920 Fifth Street
CITY: Davis
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: CG
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TELEFAX: 916-753-1510
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       46.1%;
85.7%;
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                                                        TYPE: PRT ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AMINO ACID
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Best Local Similarity
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2 DFLAEGG 8
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                                                                                                                                                                                                                                                                                                   1 DYMIOGG
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                                                                                                                     US-09-846-350A-1
SEQ ID NO 1
LENGTH: 12
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5433940-25
; Patent No. 5433940
; TILLE OF INVENTION: INHIBITORS OF THROMBIN
; TILLE OF INVENTION: INHIBITORS OF THROMBIN
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/834,259
; FILING DATE: 17-AUG-1990
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 548,388
; FILING DATE: 06-JUL-1989
; FILING DATE: 06-JUL-1989
; FILING DATE: 18-AUG-1989
                                                                                                                                                  RESULT 10
5196404-18
; Patent No. 5196404
; Patent No. 5196404
; Patent No. 5196404
; TITLE OF INVENTION: INHIBITORS OF THROMBIN
; VUMBER OF SEQUENCES: 18
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/549,388
; PILING DATE: 06-JUL-1990
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 395,482
; RILING DATE: 18-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ٠;
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Pred. No. 2e+02;
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Best Local Similarity 42.9
Matches 3; Conservative
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Matches 3; Conserv
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2 DFLAEGG 8
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LENGTH: 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 95616
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 720 KB storage
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Macintosh 6.0.7
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MEDIUM TYPE: Diskette, 3.50 inch, 720 KB storage COMPUTER: Apple Macintosh OPERATING SYSTEM: Macintosh 6.0.7
SOFTWARE: MicrosoftWord 4.0
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US-07-824-247-19
; Sequence 19, Application US/07824247
; Patent No. 5512482
; GENERAL INFORMATION:
; TITLE OF INVENTION: Plant Thioesterases
; VUMBER OF SEQUENCES: 48
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Calgene, Inc.
; STREET: 1920 Fifth Street
; CITY: Davis
; STATE: California
; CUNTRY: USA
; ZIP: 95616
                                                                                                   APPLICANT: Davies, Huw Maelor
APPLICANT: Pollard, Michael Roman
APPLICANT: Voekler, Toni Alois
APPLICANT: Thompson, Gregory A.
TITLE OF INVENTION: Plant Thioesterases
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
ADDRESSEE: Calgene, Inc.
STREET: 1920 Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWALING SISIEM: MACLILLOSH 6.0.7

SOFTWARE: MICTOSOFTWORD 4.0

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/662,007B

FILING DATE: 19910408

CLASSIFICATION: 435

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/514,030
FILING DATE: 26-APR-1990
APPLICATION NUMBER: 07/620,426
FILING DATE: 30-NOV-1990
ATTORNEY/AGENT INFORMATION:
NAME: Elizabeth Lassen
REGISTRATION NUMBER: 31,845
REGISTR
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Best Local Similarity 85.77
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AMINO ACID
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STATE: California
COUNTRY: USA
                                                             GENERAL INFORMATION:
APPLICANT: Davies,
APPLICANT: Pollard
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Patent No. 5344771
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Sequence 9, Appli
Sequence 2, Appli
Sequence 257, App
Sequence 18, Appl
Sequence 10815, A
Sequence 10911, A
Sequence 10911, A
Sequence 10911, A
Sequence 103, App
Sequence 11, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 1, Appli
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                                                                                                                                                                                   October 21, 2004, 07:15:25; Search time 61.75 Seconds (without alignments) 47.187 Million cell updates/sec
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| cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
| cgn2_6/ptodata/1/pubpaa/BCT_NEW_PUB.pep:*
| cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
| cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
| cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
| cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
| cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
| cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
| cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
| cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
| cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
| cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
| cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd
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US-09-573-822C-257

US-10-320-231A-18

US-10-057-475B-10911

US-10-154-884B-10911

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Maximum Match 100%
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Scoring table:
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                                                                                                                                   protein
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Sequence 281, App Sequence 281, App Sequence 878, App Sequence 881, App Sequence 881, App Sequence 882, App Sequence 883, App Sequence 546, App Sequence 583, App Sequence 583, App Sequence 583, App Sequence 55, Appl Sequence 27, Appl Sequence 27, Appl Sequence 891, App Sequence 891, App Sequence 891, Appl Sequence 892, Appl Sequence 891, Appl Sequence 891, Appl Sequence 892, Appl Sequence 894, Appl Sequence 894, Appl Sequence 895, Appl Sequence 896, Appl Sequence 897, Appl Sequence 896, Appl Sequence 896, Appl Sequence 897, Appl Sequence 896, Appl Sequence 897, Appl Sequence 897, Appl Sequence 896, Appl Sequence 897, Appl Sequence 896, Appl Sequence 897, Appl Sequence 8
           5 US-10-182-252A-281

US-10-182-252A-878

US-10-182-252A-878

US-10-182-252A-879

US-10-182-252A-880

US-10-182-252A-881

US-10-182-252A-881

US-10-182-252A-881

US-09-791-378-497

US-09-791-378-546

US-09-791-378-546

US-09-791-377-497

US-09-791-377-497

US-09-791-377-581

US-10-182-252A-891

US-10-182-252A-895

US-10-182-252A-895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ALIGNMENTS
US-10-788-016-9
```

Sequence 9, Application US/10788016
; Publication No. US20040141992A1
; GENERAL INFORMATION:
APPLICANT: ITCH, Kyogo
TITLE OF INVENTION: Desensitizers
; FILE REFERENCE: 3190-049
; CURRENT APPLICATION NUMBER: US/10/788,016
; CURRENT FILING DATE: 2004-02-26
; PRIOR APPLICATION NUMBER: JP P2001-260046
; PRIOR FILING DATE: 2001-08-29
; NUMBER OF SEQ ID NOS: 9
; SOFTWARE: Patentin version 3.2 ORGANISM: Artificial SEQ ID NO 9 LENGIH: 9

; OTHER INFORMATION: Designed peptide based on the peptide consisting of 9 amino acid; OTHER INFORMATION: residues from the 91st residue to the 99th residue of; OTHER INFORMATION: cyclophillin B
US-10-788-016-9 Query Match
100.0%; Score 51; DB 16;
Best Local Similarity 100.0%; Pred. No. 1.2e+06;
Matches 9; Conservative 0; Mismatches 0;

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Gaps

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Indels

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RESULT 2 US-10-447-161-85

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FEATURE: OTHER INFORMATION: Sequence located in MG024 at 351-360 and may interact with Sequence OTHER INFORMATION: in this patent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Compositions and Methods for the Detection, Diagnosis and Therapy TITLE OF INVENTION: Hematological Malignancies FILE REFERENCE: 014058-014402US CURRENT APPLICATION NUMBER: US/10/057,475B CURRENT FILING DATE: 2002-01-22
                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 5
US-10-320-231A-18
; Sequence 18, Application US/10320231A
; Publication No. US20030194405A1
; GENERAL INFORMATION:
; APPLICANT: Takeuchi, Toshihiko
; APPLICANT: Takeuchi, Toshihiko
; TITLE OF INVENTION: Artibody Inhibiting Stem Cell Factor Activity And Use For
; TITLE OF INVENTION: Treatment Of Asthma
; TITLE OF INVENTION: Treatment Of Asthma
; FILE REFERENCE: 7430*163
; CURRENT APPLICATION NUMBER: US/10/320,231A
; CURRENT FILING DATE: 2002-12-19
; PRIOR APPLICATION NUMBER: US 60/342,174
; PRIOR FILING DATE: 2001-12-17
; NUMBER OF SEQ ID NOS: 85
; SOFTWARE: Patentin version 3.2
; SEQ ID NO 18
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                                                                                                                                                                                                                                           51.0%; Score 26; DB 10;
50.0%; Pred. No. 3.7e+02;
iive 3; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10856, Application US/10057475B Publication No. US20040002068A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: synthetic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mannion, Jane
Clapper, Jonathan David
Wang, Aijun
Ordonez, Nadia
Carter, Lauren
McNeill, Patricia Dianne
Corixa Corporation
NUMBER OF SEQ ID NOS: 804
SOFTWARE: ProtPatent version 1.0
SEQ ID NO 257
LENGTH: 10
                                                                                        TYPE: PRT ORGANISM: mycoplasma genitalium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 51.0%;
Best Local Similarity 50.0%;
Matches 4; Conservative
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                                                                                                                                                                                                                                                                                       Conservative
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1 NYLIKDGD 8
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ORGANISM: Artificial
                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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APPLICANT:
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; APPLICANT: Proteom Ltd
; TITLE OF INVENTION: Complementary peptide ligands generated from microbial genome seq; FILE REFERENCE: Microbe patent; CURRENT APPLICATION NUMBER: US/09/573,822C; CURRENT FILING DATE: 2000-05-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ) OTHER INFORMATION: Peptide consisting of 9 amino acid residues from the 91st residue ) CTHER INFORMATION: to the 99th residue of cyclophillin B US-10-788-016-2
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                                                          APPLICANT: Wang, Rong-fu TITLE OF INVENTION: Mutant Fibronectin and Tumor Metastasis FILE REFERENCE: HO-P02484US1 CURRENT APPLICATION NUMBER: US/10/447,161 CURRENT FILING DATE: 2003-05-28 PRIOR APPLICATION NUMBER: 60/383,530 PRIOR FILING DATE: 2002-05-28 NUMBER OF SEQ ID NOS: 148 SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    92.2%; Score 47; DB 16; Length 9; 88.9%; Pred. No. 1.2e+06; iive 1; Mismatches 0; Indels
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Pred. No. 1.2e+06;
1; Mismatches 0
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; TILE OF INVENTION: Desensitizers
; FILE REFERENCE: 3190-049
; CURRENT APPLICATION NUMBER: US/10/788,016
; CURRENT FILING DATE: 2004-02-26
; PRIOR APPLICATION NUMBER: PCT/JP02/08641
; PRIOR FILING DATE: 2002-08-28
; PRIOR FILING DATE: 2001-08-29
; PRIOR FILING DATE: 2001-08-29
; NUMBER OF SEQ ID NOS: 9
; SOFTWARE: Patentin version 3.2
; LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Synthetic Peptide US-10-447-161-85
Sequence 85, Application US/10447161
Publication No. US20040023314A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                      92.2%;
88.9%;
                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local Similarity 88.9%
...hag 8; Conservative
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Best Local Similarity 88.9
Matches 8; Conservative
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DFMIQGGDF 9
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DFMIQGGDF 9
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ORGANISM: Artificial
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us-09-720-469a-40.rapb

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APPLICANT: Gargete, Alexander
APPLICANT: Gargete, Alexander
APPLICANT: Manion, June
APPLICANT: Corixa Corporation
TITLE OF INVENTION: Compositions and Methods for the Detection, Diagnosis and Therapy
TITLE OF INVENTION: Onload: Jul 1521US
CURRENT FILING DATE: 2002-05-23
FRICA PAPLICATION NUMBER: US 60/186,126
FRICA PAPLICATION NUMBER: US 60/180,479
FRICA PAPLICATION NUMBER: US 60/200,545
FRICA RAPLICATION NUMBER: US 60/200,545
FRICA RAPLICATION NUMBER: US 60/200,303
FRICA RAPLICATION NUMBER: US 60/200,203
FRICA RAPLICATION NUMBER: US 60/200,303
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PRIOR APPLICATION NUMBER: US 60/206,201
PRIOR FILING DATE: 2000-05-22
PRIOR APPLICATION NUMBER: US 60/218,950
PRIOR FILING DATE: 2000-07-14
PRIOR PILING DATE: 2000-07-14
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 2000-08-03
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 10979
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 10911
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
; OTHER INFORMATION: Ly1484 short MHC HLA A2 class I binding peptide
US-10-057-475B-10911
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; OTHER INFORMATION: Ly1484 long MHC HLA A2 class I binding peptide US-10-154-884B-10856
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 66.7
Matches 4; Conservative
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2 LQGGSF 7
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US-10-154-884B-10856
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APPLICANT: Clapper, Jonathan David
APPLICANT: Clapper, Jonathan David
APPLICANT: Wang, Aijun
APPLICANT: Ordonez, Nadia
APPLICANT: Carter, Lauren
APPLICANT: Corixa Corporation
TITLE OF INVENTION: Compositions and Methods for the Detection, Diagnosis and Therapy
TITLE OF INVENTION: Hematological Malignancies
FILE REFERENCE: 014058-014402US
CURRENT APPLICATION NUMBER: US/10/057,475B
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NUMBER OF SEQ ID NOS: 10979
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 10856
LENGTH: 9
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                                        PRIOR FILING DATE: 2000-03-01
PRIOR FILING DATE: 2000-03-07
PRIOR FILING DATE: 2000-03-17
PRIOR FILING DATE: 2000-04-27
PRIOR PILING DATE: 2000-04-28
PRIOR PILING DATE: 2000-04-28
PRIOR FILING DATE: 2000-05-01
PRIOR FILING DATE: 2000-05-01
PRIOR FILING DATE: 2000-05-01
PRIOR FILING DATE: 2000-05-01
PRIOR FILING DATE: 2000-05-04
PRIOR FILING DATE: 2000-05-22
PRIOR FILING DATE: 2000-07-14
PRIOR FILING DATE: 2000-07-14
PRIOR FILING DATE: 2000-07-14
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PRIOR APPLICATION NUMBER: US 60/186,126
PRIOR FILING DATE: 2000-03-01
PRIOR APPLICATION NUMBER: US 60/190,479
PRIOR FILING DATE: 2000-03-17
PRIOR APPLICATION NUMBER: US 60/200,545
PRIOR FILING DATE: 2000-04-27
PRIOR FILING DATE: 2000-04-27
PRIOR APPLICATION NUMBER: US 60/200,303
PRIOR FILING DATE: 2000-04-28
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FILING DATE: 2000-05-01
APPLICATION NUMBER: US 60/202,084
FILING DATE: 2000-05-04
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ORGANISM: Artificial Sequence
FEATURE:
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Best Local Similarity 66.7
Matches 4; Conservative
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APPLICANT: FORSGAARD, ANDERS
APPLICANT: BUNGK, SOREN
APPLICANT: BRUNAK, SOREN
APPLICANT: BUNS, SOREN
APPLICANT: BUNS, SOREN
APPLICANT: BUNS, SOREN
APPLICANT: CORBET, SYLVIE
APPLICANT: CORBET, SYLVIE
APPLICANT: CORBET, SYLVIE
APPLICANT: HANSEN, JAN
TITLE OF INVENTION: COUTROL OF HIV INFECTIONS
TITLE OF INVENTION: COOTROL OF HIV INFECTIONS
TITLE OF INVENTION: COOTROL OF HIV INFECTIONS
TITLE OF INVENTION: HIV PEPTIDE AND NUCLEIC ACIDS ENCODING THEM FOR DIAGNOSIS AND
TITLE OF INVENTION: COOTROL OF HIV INFECTIONS
TITLE OF INVENTION NUMBER: US/10/182,252A
CURRENT APPLICATION NUMBER: US 00610017.6
PRIOR FILING DATE: 2000-01-29
PRIOR FILING DATE: 2000-01-31
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 1388
SOFTWARE: PATENTIN VET: 2.1
SEQ ID NO 872
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Patent No. US20020161188A1
GENERAL INFORMATION:
APPLICANT: Jackowski, George
TITLE OF INVENTION: Biopolymer Marker Indicative of Disease State Having a Molecular V
TITLE OF INVENTION: of 1020 Daltons
FILE REFERENCE: 2132.025
CURRENT APPLICATION NUMBER: US/09/846,350
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US-10-182-252A-872
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                    CURRENT APPLICATION NUMBER: US/10/363,791
CURRENT FILING DATE: 2003-03-07
PRIOR APPLICATION NUMBER: JP 2000-274218
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 227
SOFTWARE: Patentin version 3.2
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Best Local Similarity 55.6%;
Matches 5; Conservative (
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; ORGANISM: Homo sapiens
US-10-363-791-103
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                                                                                                                                                                                                                     SEQ ID NO 103
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Sequence 10911, Application US/10154884B

Publication No. US20040005561A1

Sequence 10911, Application US/10154884B

Publication No. US20040005561A1

APPLICANT: Algate, Paul A.

APPLICANT: Corixa Corporation

ITILE OF INVENTION: Compositions and Methods for the Detection, Diagnosis and Therapy

FILE OF INVENTION: Compositions and Methods for the Detection, Diagnosis and Therapy

FILE OF INVENTION: Compositions and Methods for the Detection, Diagnosis and Therapy

FILE OF INVENTION: Hematological Malignancies

CURRENT APPLICATION NUMBER: US 60/186,126

PRIOR PLING DATE: 2000-03-21

PRIOR PLING DATE: 2000-04-27

PRIOR APPLICATION NUMBER: US 60/200,303

PRIOR APPLICATION NUMBER: US 60/200,303

PRIOR PLING DATE: 2000-04-28

PRIOR PLING DATE: 2000-05-04-28

PRIOR PLING DATE: 2000-05-05-05

PRIOR PLING DATE: 2000-05-05

PRIOR PLING DATE: 2000-05-05-05

PRIOR PLI
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; Bublication No. US20040029197A1
; GENERAL INFORMATION:
; APPLICANT: TAKIMOTO, Masato
; APPLICANT: KUZUMAKI, No. US20040029197Aloru
; APPLICANT: SATO, No. US20040029197Aliyuki
; APPLICANT: SAHARA, Hiroeki
; TITLE OF INVENTION: A novel human cancer/testis-associated gene thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                   Gaps
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  Pred. No. 1.2e+06;
1; Mismatches 1; Indels
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66.7%;
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Best Local Similarity 66.7
Matches 4; Conservative
Best Local Similarity 66.7
Matches 4; Conservative
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LQGGSF 7
                                                                                               4 IOGGDF 9
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LQGGSF 7
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US-10-363-791-103
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LENGTH: 9
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TYPE: PRT
CRGANISM: Homo sapiens
US-09-845-725-1
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6 FMVRGG 11
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                                                                        RESULT 15
US-09-845-725-1
                                                                                                                                                                                                                                                                                                                      SEQ ID NO 1
LENGTH: 12
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Patent No. US20020160422A1
GENERAL INFORMATION:
APPLICANT: Jackowski, George
TITLE OF INVENTION: BIOPOLYMER MARKER INDICATIVE OF DISEASE STATE HAVING A MOLECULAR TITLE OF INVENTION: 0F 1077 DALTONS
FILE REFERENCE: 2132.026
CURRENT FILING DATE: 2001-04-30
NUMBER OF SEQ ID NOS: 1
SOFTWARE: Patentin version 3.1
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Pred. No. 9.5e+02;
3; Mismatches 1; Indels
                                                                                                                                                                     Query Match 47.1%; Score 24; DB 9; Length 10; Best Local Similarity 42.9%; Pred. No. 8.6e+02; Matches 3; Conservative 3; Mismatches 1; Indels
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US-10-356-257-161
; Sequence 161, Application US/10356257
; Publication No. US20040087767A1
; GENERAL INFORMATION:
; APPLICANT: LAZARUS, ROBERT A.
; APPLICANT: MAUN, HENRY R.
; TITLE OF INVENTION: FVIIA ANTAGONISTS
; FILE REFERENCE: P1950R1
; CURRENT APPLICATION NUMBER: US/10/356,257
; CURRENT FILING DATE: 2003-01-30
; PRIOR FILING DATE: 2003-01-30
; PRIOR FILING DATE: 2002-02-06
; NUMBER OF SEQ ID NOS: 355
; LENGTH: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: sequence is synthesized US-10-356-257-161
      CURRENT FILING DATE: 2001-04-30 NUMBER OF SEQ ID NOS: 1 SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial sequence
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Best Local Similarity 42.9
Matches 3, Conservative
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ORGANISM: Homo sapiens
                                                         SEQ ID NO 1
LENGTH: 10
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                    1 DYMIQGG 7
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1 DFLAEGG 7
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2 DFLAEGG 8
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LENGTH: 11
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Sequence 1. Application US/09845725

Sequence 1. Application US/09845725

Sequence 1. Application US/09845725

Sequence 1. US20020161185A1

GENERAL INFORMATION:

APPLICANT: Jackowski, George

TITLE OF INVENTION: OF 1206 DALTONS

FILE REFERENCE:

CURRENT FILING DATE: 2001-04-30

NUMBER OF SEQ ID NOS: 1

SOFTWARE: Patentin version 3.1
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

October 21, 2004, 07:12:54; Search time 15.75 Seconds (without alignments) 54.981 Million cell updates/sec Run on:

US-09-720-469A-40 51 1 DYMIQGGDF 9 Title: Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

283416 seqs, 96216763 residues Searched: 1582 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 12

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

PIR 79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	Iq H2 chain - Paci	sperm-activating p	glycoprotein hormo	T-cell receptor be	kinesin light chai	phate	Ig mu chain J regi	33K protein 3218 -	hemoglobin, extrac	T-cell receptor be	cholecystokinin -	cholecystokinin -	leucokinin VIII -	e]	۲,	Б	retinal oxidase -	protein kinase C i	T-cell receptor be	T-cell receptor ga	Trimeresurus serin	recep.		: facto	yte growth	eptor g	activating	_	bradykinin-potenti
SUMMARIES	ΔΙ	B60769	E60589	A61309	PH1458	S43170	PN0165	F33932	PU0029	865730	PT0542	PQ0012	A43001	JS0318	QDRB	PT0250	PH1343	S60354	PN0044	PT0218	141946	A61575	S26544	A23751	YFPG	95	_	058	S57575	719
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ò	Query Match		٩.	39.5	9.	9.	7.	7.	7.	37.3	5	•	35.3	Ŋ.	S.	35.3	35.3	5	35.3	5	35,3	2	'n.	ო	ω.	ب	ω.	ω.	m	ъ.
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hypothetical prote calliFMRFamide 5 - calliFMRFamide 6 - calliFMRFamide 7 - polygalacturonase polygalacturonase monodehydroascorba sperm-activating p sperm-activating p sperm-activating p sperm-activating p sperm-activating p rell receptor be 1g H chain V-D-J r Z protein - guinea hypothetical prote	1993 # 100k, J rertebr 10:2123	DB 2; Length 11; 2.2e+02; ches 0; Indels 0; Gaps 0;
\$19015 E41978 F41978 G41978 C61440 A52880 A52880 A60588 C60527 E39572 D60589 D60589 S23168 T46794	ALIGNMENTS c hagfish (fragment) s stouti (Pacific hagfish) #sequence_revision 03-Jun- elt, I.M.; Gooley, A.A.; H 3-3828, 1990 H chains in a primitive v A60769; MUID:91060965; PMI tein V tein	Score 23; DB Pred. No. 2.2 1; Mismatches
4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	(f) (Pac) -rey 5990 in in UID	1. % % 0.
H 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	hagfish stouti ('equence_t, I.M.; 1828, 1916) i chains (0769; MU	45.1%; 80.0%; vative
и п п п п п п п п п п п п п п п п п п п	acific hartretus str 1993 #seq 0769 seppelt, 3823-38 ct Ig H c ber: A607 0769 minary 1 cHAN>	h Similarity 80. 4; Conservative
1	- Pac Bptatr Jun-19 July, S July, S Stinct B607 Stinct B007 Stinct Type: 1-11	ch 1 Sir 4;
0 E E E E E E E E E E E E E E E E E E E	RESULT 1 B60769 Ig H2 chain - Pacific C, Species: Eptatretus C; Date: 03-Jun-1993 #s C; Accession: B60769 R; Hanley, P.J.; Seppel J. Immunol. 145, 3823-A; Title: Distinct Ig HA; A; Reference number: A6 A; Accession: B60769 A; Status: preliminary A; Molecule type: prote A; Residues: 1-11 < HAN>	Query Match Best Local Matches

Decrees: Echinometra mathaei C;Species: Echinometra mathaei C;Species: Echinometra mathaei C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 16-Aug-2004 C;Accession: E60589 R;Yoshino, K.I.; Kajiura, H.; Nomura, K.; Takao, T.; Shimonishi, Y.; Kurita, M.; Yamaguch Comp. Biochem. Physiol. B 94, 739-751, 1989 A;Title: A halogenated amino acid-containing sperm activating peptide and its related per Otus nudus, Echinometra mathaei and Heterocentrotus mammillatus. A;Reference number: A60527 A;Reference number: A60527 A;Molecule type: protein A;Residues: 1-10 < YOS> A;Cross-references: UNIPROT:Q7M4C1 A;Note: an identical peptide was isolated from Echinometra mathaei type A and type B RESULT 2 E60589

|:||| 5 DFMIQ 9 1 DYMIQ 5

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Gaps . Query Match 39.2%; Score 20; DB 2; Length 10; Best Local Similarity 50.0%; Pred. No. 7.7e+02; Matches 3; Conservative 1; Mismatches 2; Indels

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2 YSLSGG 7 q

2 YMIQGG 7

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E) RESULT :

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A;Residues: 1-12 <CHE>
A;Cross-references: UNIPROT:Q13865; EMBL:X69658; NID:g468786; PIDN:CAA49349.1; PID:g46878
              A; Molecule type: DNA
A; Residues: 1-12 <CHE>
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glycoprotein hormones alpha chain - hamster (fragment)

N; Alternate names: choriogonadotropin alpha chain; luteinizing hormone alpha chain; lutr
C; Species: Cricetinae gen. sp. (hamster)
C; Date: 17-Jul-1994 #sequence_revision 17-Jul-1994 #text_change 09-Jul-2004
C; Accession: A61309
R; Glem, S.D.; Nahm, H.S.; Greenwald, G.S.; Ward, D.N.
Endocrinology 111, 1263-1269, 1982
A; Title: Isolation and characterization of hamster luteinizing hormone.
A; Reference number: A61309; MUID: 83003498; PMID: 6889489
A; Status: preliminary
A; Molecule type: protein
A; Residues: 1-12 < GLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       T-cell receptor beta chain (clone 332/1K) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 15-Mar-2004
C;Date: 10-Mar-1994; S26543
C;Accession: PH1458; S26543
R;Casanova, J.L.; Martinon, F.; Gournier, H.; Barra, C.; Pannetier, C.; Regnault, A.; KcJ. Exp. Med. 177, 811-820, 1993
A;Title: T cell receptor selection by and recognition of two class I major histocompatik
A;Reference number: PH1430; MUID:93171821; PMID:8436911
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Molecule type: mRNA
A;Residues: 1-12 <CAS>
A;Experimental source: cytolytic T-lymphocyte, clone 332/1K
R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Barra, C.; Wid
J. Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T cell receptor
A;Reference number: S26512; MUID:92364546; PMID:1380061
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843170
kinesin light chain - human (fragment)
C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 13-Jan-1995 #sequence_revision 18-Aug-2000 #text_change 09-Jul-2004
C; Accession: 843170
C; Accession: Prown, A.; Jones, T.A.; Sheer, D.
Submitted to the EMBL Data Library, December 1992
A; Description: Promoter first exon/intron characterization and chromosomal location A; Reference number: 843170
A; Reference number: 843170
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C;Keywords: receptor; T-cell
                                                                                                                                                                                                                                                                                                                                                                                                                                              39.2%; Score 20; DB 2; I
50.0%; Pred. No. 9.3e+02;
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C, Superfamily: glycoprotein hormones alpha chain
C, Keywords: glycoprotein
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Matches 3; Conservative
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Best Local Similarity
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DFTMOG 10
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A; Residues: 1-12 <CA2>
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QGSDY 9
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C;Species: Fusarium sporotrichioides
C;Date: 05-Aug-1994 #sequence_revision 06-Jan-1995 #text_change 09-Jul-2004
C;Accession: PN0165
R;Fukaya, N.; Chow, L.P.; Sugiura, Y.; Tsugita, A.; Ueno, Y.; Tabuchi, K.
submitted to JIPID, May 1994
A;Description: Two dimensional polyacrylamide gel electrophoresis of Fusarium sporotrichi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ig mu chain J region (E7) - mouse
C;Species: Mus musculus (house mouse)
C;Date: 09-Mar-1990 #sequence_revision 09-Mar-1990 #text_change 09-Jul-2004
C;Accession: F33932
R;Baccala, R.; Vo Quang, T.; Gilbert, M.; Ternynck, T.; Avrameas, S.
Proc. Natl. Acad. Sci. U.S.A. 86, 4624-4628, 1989
A;Title: Two murine natural polyreactive autoantibodies are encoded by nonmutated germ-lb.
A;Reference number: A33932; MUID:892823; PMID:2499887
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C, Keywords: immunoglobulin
                                                                                                                                                                                                                                                                                                                       triose-phosphate isomerase (EC 5.3.1.1) - fungus (Fusarium sporotrichioides) (fragment)
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C;Species: Oryza sativa (rice)
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 11-Apr-1995
C;Accession: PU0029
R;Tsugita, A.; Miyatake, N.
                                                                             Gaps
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                       39.2%; Score 20; DB 2; Le
60.0%; Pred. No. 9.3e+02;
tive 2; Mismatches 0;
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Jocal Similarity 75.0%; Pred. No. 1.2e+03;
Jess 3; Conservative 1; Mismatches 0;
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C, Keywords: intramolecular oxidoreductase; isomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: protein
A; Residues: 1-10 <FUK>
A; Cross-references: UNIPROT: Q7M4X7
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A; Accession: PN0165
Query Match
Best Local Similarity
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A,Molecule type: mRNA
A,Residues: 1-10 <BAC>
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C;Species: Leucophaea maderae (Madeira cockroach)
C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 09-Jul-2004
C;Accession: JS0318
R;Holman, G.M.; Cook, B.J.; Nachman, R.J.
Comp. Biochem. Physiol. C 88, 31-34, 1987
A;Title: Isolation, primary structure and synthesis of leucokinins VII and VIII: the fine A;Reference number: JS0317
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A;Cross-references: UNIPROT:P19990
C;Comment: Leucokinins, a family of cephalomyotropic peptides, stimulate contractile acti
C;Keywords: amidated carboxyl end; cephalomyotropic peptide
F;8/Modified site: amidated carboxyl end (Gly) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Macropus eugenii (tammar wallaby)
C;Date: 30-Oct-1992 #sequence_revision 30-Oct-1992 #text_change 09-Jul-2004
C;Accession: A43001; PQ0012
R;Fan, Z.W.; Eng, J.; Shaw, G.; Yalow, R.S.
Peptides 9, 429-431, 1988
A;Title: Cholecystokinin octapeptide purified from brains of Australian marsupials.
A;Reference number: PQ0012; MUID:88234141; PMID:3375140
A;Reference protein
A;Residues: 1-8 <FAN>
      C;Accession: PQ0012
R;Fan, Z.W.; Eng, J.; Shaw, G.; Yalow, R.S.
Peptides 9, 429-431, 1988
A;Title: Cholecystokinin octapeptide purified from brains of Australian marsupials.
A;Reference number: PQ0012; MUID:88234141; PMID:3375140
A;Accession: PQ0012
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                                                                                                                                                                                                                         A;Cross-references: UNIPROT:P30369
C;Superfamily: gastrin
C;Keywords: amidated carboxyl end; hormone; neuropeptide; sulfoprotein
F;2/Binding site: sulfate (Tyr) (covalent) #status predicted
F;8/Modified site: amidated carboxyl end (Phe) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPROT:P30369
C;Superfamily: gastrin
C;Keywords: amidated carboxyl end; hormone; neuropeptide; sulfoprotein
F;2/Binding site: sulfate (Tyr) (covalent) #status predicted
F;8/Modified site: amidated carboxyl end (Phe) #status predicted
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A; Molecule type: protein
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Best Local Similarity
Matches 3; Conserv
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C; Species: Lumbricus terrestris (common earthworm)
C; Date: 06-Dec-1996 #sequence_revision 13-Mar-1997 #text_change 13-Mar-1997
C; Date: 06-Dec-1996 #sequence_revision 13-Mar-1997 #text_change 13-Mar-1997
C; Accession: S65730
C; Accession: S65730
C; Accession: S65721; MUID: 96176855; PMID: 8597573
A; Reference number: S65721; MUID: 96176855; PMID: 8597573
A; Accession: S65730
A; Status: preliminary
A; Molecule type: protein
A; Residues: 1-12 < FUS>
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C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0542
R;Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hemoglobin, extracellular, component - earthworm (Lumbricus terrestris) (fragment)
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N;Alternate names: CCK
C;Species: Dasyurus viverrinus (southeastern quoll)
C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 09-Jul-2004
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Pred. No. 1.5e+03;
2; Mismatches 0; Indels
                                                                                                                                                                                                                            Score 19; DB 2; Length 11;
Pred. No. 1.3e+03;
1; Mismatches 1; Indels
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A;Molecule type: mRNA
A;Residues: 1-7 <FEE>
A;Experimental source: day 18 fetal thymus, strain BALB/c
C;Keywords: T-cell receptor
submitted to JIPID, April 1993
A; Reference number: PS0208
A; Accession: PU0029
A; Molecule type: protein
A; Residues: 1-11 <TSU>
A; Experimental source: bran
C; Comment: molecular weight 33K, pI 6.0.
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                                                                                                                                                                                                                            Query Match 37.3%;
Best Local Similarity 60.0%;
Matches 3; Conservative
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Matches 3; Conserv
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3 EGGPF 7
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C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 20-Jun-2000 #sequence_revision 20-Jun-2000 #text_change 16-Aug-2004
C;Accession: A01422
R;Monnier, M.; Dudler, L.; Gachter, R.; Maier, P.F.; Tobler, H.J.; Schoenenberger, G.A.
Experientia 33, 548-552, 1977
A;Title: The delta sleep inducing peptide (DSIP). Comparative properties of the original A;Reference number: A01422
A;Molecule type: protein
A;Molecule type: protein
A;Residues: 1-9 <MON>
A;Cross-references: UNIPROT:P01158
C;Comment: This peptide was obtained from dialysates of occipital venous sinus blood fro of recipient rabbits, it induces spindle and delta EEG activity and reduced motor activ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 15
PT0250
Ig heavy chain CRD3 region (clone 2-109B) - human (fragment)
C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C; Accession: PT0250
C; Accession: PT0250
C; Accession: PT0250
J. Exp. Med. 173, 395-407, 1991
A; Title: Preferential utilization of specific immunoglobulin heavy chain diversity and j
A; Reference number: PT0222; MUD:91108337; PMID:1899102
A; Molecule type: DNA
A; Residues: 1-11 < YAM>
A; Residues: 1-11 < YAM>
A; Residues: 1-11 < YAM>
C; Keywords: heterotetramer; immunoglobulin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  35.3%; Score 18; DB 2; Length 9; 100.0%; Pred. No. 2.8e+05; tive 0; Mismatches 0; Indels
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GenCore version 5.1.6
(c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

Run on:

October 21, 2004, 07:05:04; Search time 86 Seconds (without alignments) 60.214 Million cell updates/sec

US-09-720-469A-40 51 1 DYMIQGGDF 9 Title: Perfect score:

Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

1825181 seqs, 575374646 residues Searched: 4909 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 12

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

UniProt_02:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Q9N2B8	Q9N2B9	Q9N2C0	046712	ESTA SCHGA	ASCL_ALLAS	Q90ZV5	THYF PIG	Q7M3Č5	AMPN HELAM	Q6LC <u>I</u> 4	Q96QA7	Q7M4B7	Q8MJ52
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18	18	18	18	17.5	17	17	17	17	17	17	17	17	17
32	33	34	35	36	37	38	39	40	41	42	43	44	45

# ALIGNMENTS

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0;
                                                                                                    Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
1818 TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ul-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2002 (TrEMBLrel. 21, Last annotation update)
10-JUN-2002 (TrEMBLrel. 21, Last annotation update)
LOW density lipoprotein receptor-related protein (Fragment).
Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                        "Splice variants of rat TR4 orphan receptor: differential expression of novel sequences in the 5'-untranslated region and C-terminal
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Yoshikawa T., Makino S., Gao X.M., Xing G.Q., Chuang D.M.,
Detera-Wadleigh S.D.;
                                                                                                                                                                                                                                                                                                                                                                                   Score 23; DB 2; Length 7; Pred. No. 1.8e+06; 1; Mismatches 0; Indels
                                                     Last sequence update)
Last annotation update)
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            7 AA.
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                                      01-NOV-1996 (TrEMBLrel. 01, Created) 01-NOV-1996 (TrEMBLrel. 01, Last seq 01-JUN-2003 (TrEMBLrel. 24, Last ann TR4-NS orphan receptor (Fragment).
            PRT;
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Endocrinology 137:1562-1571(1996).
EMBL; U59125; AAB02827.1; -.
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80.0%;
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NCBI_TaxID=9031;
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MEDLINE=94010310; PubMed=8406013;
Onno M., Nakamura T., Hillova J., Hill M.;
"Identification of novel sequences in the repertoire of hypervariable "IRE17 genes from immortalized nonmalignant and malignant human
                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                       "The laying hen expresses two different low density lipoprotein
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MEDLINE=93228825; PubMed=8471161;
Onno M., Nakamura T., Mariage-Samson R., Hillova J., Hill M.;
"Human TRE17 oncogene is generated from a family of homologous polymorphic sequences by single-base changes.";
DNA Cell Biol. 12:107-118(1993).
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                                                                                                                                                                                                                           0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; X71371; CAC16150.1; -. SEQUENCE 12 AA; 1252 MW; 6E17AFDE83ADD87B CRC64;
                                                                                                                                  895EC68B5AB2CDC1 CRC64;
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Pred. No. 2.7e+03;
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                                           receptor-related proteins.";
J. Biol. Chem. 266:19079-19087(1991).
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05-JUL-2004 (TrEMBLrel. 27, Last seq
05-JUL-2004 (TrEMBLrel. 27, Last ann
Ribosomal protein (Fragment).
Name=rps16;
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                                                                                                                                  9 AA; 1039 MW;
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3; Conservative
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DYMVK 9
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MVEGAD
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Schneider W.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TRE17 protein.
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SEQUENCE
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                                                                                                                                                                                                                           Matches
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Q9H3W2
ID Q9H3W2
DT Q1-MI
DT O1-MI
DT O1-MI
DT O1-MI
DT O1-MI
DE TRE-J
GN Name
OS HOMO
OC MAMMI
OX NCBI
RN [1]
RP SEQUI
RX MEDL
RA ONDO
RT "Idea
RT "Idea
RT "Est
RT Gene
RN [2]
RP SEQUI
RX MEDL
RA ONDO
RT Gene
RN [2]
RP SEQUI
RX MEDL
RA ONDO
RT "HUM
RT FEI
RP SEQUI
RX MEDL
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RY HILL
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RY SEQUI
RY SEQUI
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270Y64
ID Q70Y
AC Q70Y
DT 05-U
DT 05-U
DT 05-U
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DT 05-C
OG Chlo
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EMBL; AJ505409; CAD45529.1; -...
GO: GO:0003735; F:structural constituent of ribosome; IEA.
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
lamiids; Lamiales; Lamiaceae; Nepetoideae; Ocimeae; Plectranthus.
NCBI_TaxID=204208;
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Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots, asterids,
lamiids, Lamiales, Lamiaceae, Nepetoideae, Ocimeae, Plectranthus.
NCBI_TaxID=204208;
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EMBL; AJ505409; CAD45529.1; -...
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Pred. No. 3.9e+03;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                           11 AA; 1317 MW; D8D98CB3AB587866 CRC64;
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Last annotation update)
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Ribosomal protein (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11 AA.
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Similarity 66.7%;
4; Conservative
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                                                                                                                 SEQUENCE FROM N.A. PubMed=15019625;
                                                                                                                                                                                                                                                                                                                                   Ribosomal protein.
NON TER 1
NON_TER 11
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Best Local Similarity
Matches 4; Conserv
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Chloroplast.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
lamids; Lamiales; Lamiaceae; Nepetoideae; Ocimeae; Platostoma.
NCBI_TaxID=204168;
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EMBL; AJ505368; CAD45489.1; -...
                                                                                                                                                                                                                                                                                                                                                                        Paton A., Springate D.A., Sudde S., Otieno D., Grayer R., Harley M.M., Willis F., Simonds M.S.J., Powell M.P., Savolainen V.; "Phylogeny and evolution of basils and allies (Ocimeae, Labiatae) based on three plastid DNA regions."; Mol. Phylogenet. Evol. 31:277-299(2004).

EMBL; AJ505368; CAD45489.1; -...
GO: GO:0003735; F:structural constituent of ribosome; IEA.
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Eukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
lamiids; Lamiales; Lamiaceae; Nepetoideae; Ocimeae; Platostoma.
NCBI_TaxID=204168;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  39.2%; Score 20; DB 2; Length 8; 50.0%; Pred. No. 1.8e+06; Live 1; Mismatches 2; Indels
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14-MAR-2004 (TrEMBLrel. 27, Last sequence update)
14-MAR-2004 (TrEMBLrel. 27, Last annotation update)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update) 05-JUL-2004 (TrEMBLrel. 27, Last annotation update) Ribosomal protein (Fragment).
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                                                                                                                 Platostoma fimbriatum.
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Best Local Similarity
3, Conserve
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                     PubMed=15019625;
                                                                                       Name=rps16;
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SEQUENCE
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NON TER
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CAD45489
ID CAD4
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
lamiids; Lamiales; Lamiaceae; Nepetoideae; Nepeteae; Clinopodium;
Clinopodium vulgare.
NCBI_TaxID=204223;
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EMBL; AJS05426; CAD45546.1; -...
                                          Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
lamiids; Lamiales; Lamiaceae; Nepetoideae; Nepeteae; Clinopodium.
NCBI_TaxID=204223;
                                                                                                                                                                                                                                                             Paton A., Springate D.A., Sudde S., Otieno D., Grayer R., Harley M.M., Willis F., Simonds M.S.J., Powell M.P., Savolainen V.; "Phylogeny and evolution of basils and allies (Ocimeae, Labiatae) based on three plastid DNA regions."; Mol. Phylogenet. Evol. 31:277-299 (2004).

EMBL; AJ505426; CAD45546.1; ---
GO; GO:0003735; F:structural constituent of ribosome; IEA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12 12
12 AA; 1353 MW; D43D1B51AF587866 CRC64;
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14-MAR-2004 (TrEMBLrel. 27, Last sequence update)
14-MAR-2004 (TrEMBLrel. 27, Last annotation update)
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05-JUL-2004 (TrEMBLrel. 27, Created)
       Clinopodium vulgare subsp. arundanum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best_Local Similarity 100.
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
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                                                                                                                                                                                                           SEQUENCE FROM N.A.
PubMed=15019625;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ribosomal protein.
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AC Q70Y
DT 05-J1

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Query Match
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Matches 3; Conserv
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                                                                                                                                                                                                                                                            Yoshino K.I., Kajiura H., Nomura K., Takao T., Shimonishi Y.,
Kurita M., Yamaguchi M., Suzuki N.;
"A halogenated amino acid-containing sperm activating peptide and its
related peptides isolated from the egg jelly of sea urchins,
Tripneustes gratilla, Pseudoboletia maculata, Strongylocentrotus
nudus, Echinometra mathaei and Heterocentrotus mammillatus.";
Comp. Biochem. Physiol. 94:739-751(1989).
PIR; E60589; E60589.
SEQUENCE 10 AA; 925 MW; 8B9856D87865B735 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE=Placenta;
MEDLINE=97101110; PubMed=8945637;
Chernajovsky Y., Brown A., Jones T.A.;
Chernajovsky Y., Brown A., Jones T.A.;
"Promoter first exon/intron characterization and chromosomal location of the human light chain (beta) kinesin gene.";
DNA Cell Biol. 15:965-974(1996).
EMBL; X69658; CAA49349.1; -.
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                    01-MAR-2004 (TrEMBLrel. 26, Created)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Sperm-activating peptide (Tyr-2, Ser-3,5, Ala-8, Asp-10 SAP-I).
Echinometra mathaei (Rock boring urchin).
Eukaryota; Metazoa; Echinodermata; Eleutherozoa; Echinozoa; Echinoidea; Echinoidea; Echinoidea; Echinoidea; Echinoidea; Echinometridae;
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01-NOV-1996 (TrEMBLrel. 01,
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MLRGG 5
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Eukaryota; Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia; Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Rattus.
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Glycoprotein hormones alpha chain (Fragment).
Cricetidae sp. (Hamster).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Cricetinae.
                                                                                                                                                                                                                        MEDIINE=83003498; PubMed=6889489; Glenn S.D., Nahm H.S., Greenwald G.S., Ward D.N.; "Isolation and characterization of hamster luteinizing hormone."; Endocrinology 111:1263-1269(1982).
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MEDLINE=96299786; PubMed=8661150;
Yoshikawa T., DuPont B.R., Leach R.J., Detera-Wadleigh S.D.;
"New variants of the human and rat nuclear hormone receptor, TR4: expression and chromosomal localization of the human gene.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .;
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of novel sequences in the 5'-untranslated region and C-terminal
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MEDLINE=96198747; PubMed=8612486;
Yoshikawa T., Makino S., Gao X.M., Xing G.Q., Chuang D.M.,
Detera-Wadleigh S.D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 12;
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12 AA; 1280 MW; 94A63728AA81A9CA CRC64;
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Last annotation update)
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Similarity 80.0%;
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EMBL; U59454; AAB91433.1;
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01-MAR-2002 (TrEMBLrel. 20, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
AMPA receptor subunit 2 alpha (Fragment).
Name=GluR2a;
Oreochromis mossambicus (Mozambique tilapia) (Tilapia mossambica).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostemi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Labroidei; Cichlidae; Oreochromis.
NCBI TaxID=8127;
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PIR; PN0165; PN0165.

GO; GO:0004807; F:triose-phosphate isomerase activity; IEA.

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NON_TER 10 10

SEQÜENCE 10 AA; 1142 MW; 590E4D7862C9C9C73 CPC64.
    01-MAR-2004 (TrEMBLrel. 26, Created)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Triose-phosphate isomerase (EC 5.3.1.1) (Fragment).
Fusarium sporotrichioides.
Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycetes;
Hypocreomycetidae; Hypocreales; mitosporic Hypocreales; Fusarium.
NCBI TaxID=5514;
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.10 Aa; 1153 MW; 721A6E5AAEB866C4 CRC64;
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Best Local Similarity 57.1%;
Matches 4; Conservative
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